



8

SEQUENCE LISTING

<110> Ruvkun, Gary
Ogg, Scott

<120> THERAPEUTIC AND DIAGNOSTIC TOOLS FOR
IMPAIRED GLUCOSE TOLERANCE CONDITIONS

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<140> 09/205,658
<141> 1998-12-03

<150> 08/857,076
<151> 1997-05-15

<150> 08/888,534
<151> 1997-07-07

<150> US98/10080
<151> 1998-05-15

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 Lys Asp Leu Ala Gly Gln Arg Met Val Asn Cys Thr Val Val Glu Gly
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 85 90 95
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 Thr Glu Gly Leu Val Asp Leu Arg Lys Ile Phe Pro Asn Leu Arg Val
 115 120 125
 Ile Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg
 130 135 140
 Asn Pro Asp Leu Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn
 145 150 155 160
 Gly Gly Val Arg Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr
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 180 185 190
 Asp Asn Ala Ala Glu Tyr Ala Val Thr Glu Thr Gly Leu Met Cys Pro
 195 200 205
 Arg Gly Ala Cys Glu Glu Asp Lys Gly Glu Ser Lys Cys His Tyr Leu
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 Glu Glu Lys Asn Gln Glu Gln Gly Val Glu Arg Val Gln Ser Cys Trp
 225 230 235 240
 Ser Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro
 245 250 255
 Thr Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His
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 275 280 285
 His Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys
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 Asp Ala His Leu Tyr Leu Leu Leu Gln Arg Arg Cys Val Thr Arg Glu

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Pro	Asn	Leu	Lys	Lys	Leu	Phe	Asp	Ser	Thr	Thr	Asp	Leu	Thr	Leu	Asp	
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Gln	Ser	Glu	Gly	Thr	Asn	Gly	Glu	Lys	Ala	Ile	Cys	Glu	Asp	Met	Ala	
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Met	Asp	Ile	Gly	Pro	Arg	Glu	Arg	Ile	Arg	Pro	Asn	Thr	Leu	Tyr	Ala	
610	615	620														
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625	630	635	640													
Val	Ser	Lys	Ile	Gly	Phe	Val	Arg	Thr	Ser	Tyr	Tyr	Thr	Pro	Asp	Pro	
645	650	655														
Pro	Thr	Leu	Ala	Leu	Ala	Gln	Val	Asp	Ser	Asp	Ala	Ile	His	Ile	Thr	
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675	680	685														
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690	695	700														
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Gln Gln Ala Ala Thr Ala Ala Ala Ala Ala Ala Ala Leu Gln Gln		
1635	1640	1645
Gln Gln Asn Gly Gly Arg Gly Asp Arg Leu Thr Gln Leu Pro Gly Thr		
1650	1655	1660
Gly His Leu Gln Ser Thr Arg Gly Gly Gln Asp Gly Asp Tyr Ile Glu		
1665	1670	1675
Thr Glu Pro Lys Asn Tyr Arg Asn Asn Gly Ser Pro Ser Arg Asn Gly		
1685	1690	1695
Asn Ser Arg Asp Ile Phe Asn Gly Arg Ser Ala Phe Gly Glu Asn Glu		
1700	1705	1710
His Leu Ile Glu Asp Asn Glu His His Pro Leu Val		
1715	1720	

<210> 13
<211> 139
<212> PRT
<213> *Caenorhabditis elegans*

<400> 13

Thr	Ser	Gly	Ser	Gly	Met	Gly	Pro	Thr	Thr	Leu	His	Lys	Leu	Thr	Ile
1					5				10					15	
Gly	Gly	Gln	Ile	Arg	Leu	Thr	Gly	Arg	Val	Gly	Ser	Gly	Arg	Phe	Gly
					20				25				30		
Asn	Val	Ser	Arg	Gly	Asp	Tyr	Arg	Gly	Glu	Ala	Val	Ala	Val	Lys	Val
					35				40			45			
Phe	Asn	Ala	Leu	Asp	Glu	Pro	Ala	Phe	His	Lys	Glu	Thr	Glu	Ile	Phe
					50				55		60				
Glu	Thr	Arg	Met	Leu	Arg	His	Pro	Asn	Val	Leu	Arg	Tyr	Ile	Gly	Ser
					65				70		75		80		
Asp	Arg	Val	Asp	Thr	Gly	Phe	Val	Thr	Glu	Leu	Trp	Leu	Val	Thr	Glu
					85				90			95			
Tyr	His	Pro	Ser	Gly	Ser	Leu	His	Asp	Phe	Leu	Leu	Glu	Asn	Thr	Val
					100				105			110			
Asn	Ile	Glu	Thr	Tyr	Tyr	Asn	Leu	Met	Arg	Ser	Thr	Ala	Ser	Gly	Leu
					115				120			125			
Ala	Phe	Leu	His	Asn	Gln	Ile	Gly	Gly	Ser	Lys					
					130				135						

<210> 14
<211> 62
<212> PRT
<213> *Caenorhabditis elegans*

<400> 14

Glu	Asp	Ala	Ala	Ser	Asp	Ile	Ile	Ala	Asn	Glu	Asn	Tyr	Lys	Cys	Gly
1					5				10			15			
Thr	Val	Arg	Tyr	Leu	Ala	Pro	Glu	Ile	Leu	Asn	Ser	Thr	Met	Gln	Phe
					20				25			30			
Thr	Val	Phe	Glu	Ser	Tyr	Gln	Cys	Ala	Asp	Val	Tyr	Ser	Phe	Ser	Leu
					35				40		45				
Val	Met	Trp	Glu	Thr	Leu	Cys	Arg	Cys	Glu	Asp	Gly	Asp	Val		
					50				55		60				

<210> 15
<211> 31
<212> PRT
<213> *Caenorhabditis elegans*

<400> 15

Lys	Pro	Ala	Met	Ala	His	Arg	Asp	Ile	Lys	Ser	Lys	Asn	Ile	Met	Val
1					5				10			15			
Lys	Asn	Asp	Leu	Thr	Cys	Ala	Ile	Gly	Asp	Leu	Gly	Leu	Ser	Leu	
					20				25			30			

<210> 16
<211> 72
<212> PRT
<213> *Caenorhabditis elegans*

<400> 16
Ile Pro Tyr Ile Glu Trp Thr Asp Arg Asp Pro Gln Asp Ala Gln Met
1 5 10 15
Phe Asp Val Val Cys Thr Arg Arg Leu Arg Pro Thr Glu Asn Pro Leu
20 25 30
Trp Lys Asp His Pro Glu Met Lys His Ile Met Glu Ile Ile Lys Thr
35 40 45
Cys Trp Asn Gly Asn Pro Ser Ala Arg Phe Thr Ser Tyr Ile Cys Arg
50 55 60
Lys Arg Met Asp Glu Arg Gln Gln
65 70

<210> 17
<211> 150
<212> PRT
<213> *Caenorhabditis elegans*

<400> 17
Tyr Phe Glu Ser Val Asp Arg Phe Leu Tyr Ser Cys Val Gly Tyr Ser
1 5 10 15
Val Ala Thr Tyr Ile Met Gly Ile Lys Asp Arg His Ser Asp Asn Leu
20 25 30
Met Leu Thr Glu Asp Gly Lys Tyr Val His Ile Asp Phe Gly His Ile
35 40 45
Leu Gly His Gly Lys Thr Lys Leu Gly Ile Gln Arg Asp Arg Gln Pro
50 55 60
Phe Ile Leu Thr Glu His Phe Met Thr Val Ile Arg Ser Gly Lys Ser
65 70 75 80
Val Asp Gly Asn Ser His Glu Leu Gln Lys Phe Lys Thr Leu Cys Val
85 90 95
Glu Ala Tyr Glu Val Met Trp Asn Asn Arg Asp Leu Phe Val Ser Leu
100 105 110
Phe Thr Leu Met Leu Gly Met Glu Leu Pro Glu Leu Ser Thr Lys Ala
115 120 125
Asp Leu Asp His Leu Lys Lys Thr Leu Phe Cys Asn Gly Glu Ser Lys
130 135 140
Glu Glu Ala Arg Lys Phe
145 150

<210> 18
<211> 113
<212> PRT
<213> *Caenorhabditis elegans*

<400> 18
Ser Pro Leu Asp Pro Val Tyr Lys Leu Gly Glu Met Ile Ile Asp Lys
1 5 10 15
Ala Ile Val Leu Gly Ser Ala Lys Arg Pro Leu Met Leu His Trp Lys
20 25 30
Asn Lys Asn Pro Lys Ser Asp Leu His Leu Pro Phe Cys Ala Met Ile
35 40 45
Phe Lys Asn Gly Asp Asp Leu Arg Gln Asp Met Leu Val Leu Gln Val
50 55 60
Leu Glu Val Met Asp Asn Ile Trp Lys Ala Ala Asn Ile Asp Cys Cys
65 70 75 80
Leu Asn Pro Tyr Ala Val Leu Pro Met Gly Glu Met Ile Gly Ile Ile
85 90 95

Glu Val Val Pro Asn Cys Lys Thr Ile Phe Glu Ile Gln Val Gly Thr
100 105 110
Gly

<210> 19
<211> 106
<212> PRT
<213> *Caenorhabditis elegans*

<400> 19
Leu Ala Phe Val Trp Thr Asp Arg Glu Asn Phe Ser Glu Leu Tyr Val
1 5 10 15
Met Leu Glu Lys Trp Lys Pro Pro Ser Val Ala Ala Ala Leu Thr Leu
20 25 30
Leu Gly Lys Arg Cys Thr Asp Arg Val Ile Arg Lys Phe Ala Val Glu
35 40 45
Lys Leu Asn Glu Gln Leu Ser Pro Val Thr Phe His Leu Phe Ile Leu
50 55 60
Pro Leu Ile Gln Ala Leu Lys Tyr Glu Pro Arg Ala Gln Ser Glu Val
65 70 75 80
Gly Met Met Leu Leu Thr Arg Ala Leu Cys Asp Tyr Arg Ile Gly His
85 90 95
Arg Leu Phe Trp Leu Leu Arg Ala Glu Ile
100 105

<210> 20
<211> 139
<212> PRT
<213> *Caenorhabditis elegans*

<400> 20
Glu Tyr Trp Ile Val Thr Glu Phe His Glu Arg Leu Ser Leu Tyr Glu
1 5 10 15
Leu Leu Lys Asn Asn Val Ile Ser Ile Thr Ser Ala Asn Arg Ile Ile
20 25 30
Met Ser Met Ile Asp Gly Leu Gln Phe Leu His Asp Asp Arg Pro Tyr
35 40 45
Phe Phe Gly His Pro Lys Lys Pro Ile Ile His Arg Asp Ile Lys Ser
50 55 60
Lys Asn Ile Leu Val Lys Ser Asp Met Thr Thr Cys Ile Ala Asp Phe
65 70 75 80
Gly Leu Ala Arg Ile Tyr Ser Tyr Asp Ile Glu Gln Ser Asp Leu Leu
85 90 95
Gly Gln Val Gly Thr Lys Arg Tyr Met Ser Pro Glu Met Leu Glu Gly
100 105 110
Ala Thr Glu Phe Thr Pro Thr Ala Phe Lys Ala Met Asp Val Tyr Ser
115 120 125
Met Gly Leu Val Met Trp Glu Val Ile Ser Arg
130 135

<210> 21
<211> 61
<212> PRT
<213> *Caenorhabditis elegans*

<400> 21
Ile Gly Phe Asp Pro Thr Ile Gly Arg Met Arg Asn Tyr Val Val Ser
1 5 10 15
Lys Lys Glu Arg Pro Gln Trp Arg Asp Glu Ile Ile Lys His Glu Tyr
20 25 30
Met Ser Leu Leu Lys Lys Val Thr Glu Glu Met Trp Asp Pro Glu Ala
35 40 45
Cys Ala Arg Ile Thr Ala Gly Cys Ala Phe Ala Arg Val
50 55 60

<210> 22
<211> 20
<212> PRT
<213> *Caenorhabditis elegans*

<400> 22
Pro Ile Thr Asp Phe Gln Leu Ile Ser Lys Gly Arg Phe Gly Lys Val
1 5 10 15
Phe Lys Ala Gln
20

<210> 23
<211> 163
<212> PRT
<213> *Caenorhabditis elegans*

<400> 23
Thr Asp Ser Glu Thr Arg Ser Arg Phe Ser Leu Gly Trp Tyr Asn Asn
1 5 10 15
Pro Asn Arg Ser Pro Gln Thr Ala Glu Val Arg Gly Leu Ile Gly Lys
20 25 30
Gly Val Arg Phe Tyr Leu Leu Ala Gly Glu Val Tyr Val Glu Asn Leu
35 40 45
Cys Asn Ile Pro Val Phe Val Gln Ser Ile Gly Ala Asn Met Lys Asn
50 55 60
Gly Phe Gln Leu Asn Thr Val Ser Lys Leu Pro Pro Thr Gly Thr Met
65 70 75 80
Lys Val Phe Asp Met Arg Leu Phe Ser Lys Gln Leu Arg Thr Ala Ala
85 90 95
Glu Lys Thr Tyr Gln Asp Val Tyr Cys Leu Ser Arg Met Cys Thr Val
100 105 110
Arg Val Ser Phe Cys Lys Gly Trp Gly Glu His Tyr Arg Arg Ser Thr
115 120 125
Val Leu Arg Ser Pro Val Trp Phe Gln Ala His Leu Asn Asn Pro Met
130 135 140
His Trp Val Asp Ser Val Leu Thr Cys Met Gly Ala Pro Pro Arg Ile
145 150 155 160
Cys Ser Ser

<210> 24
<211> 44
<212> PRT
<213> *Caenorhabditis elegans*

<400> 24

Arg Ala Phe Arg Phe Pro Val Ile Arg Tyr Glu Ser Gln Val Lys Ser
1 5 10 15
Ile Leu Thr Cys Arg His Ala Phe Asn Ser His Ser Arg Asn Val Cys
20 25 30
Leu Asn Pro Tyr His Tyr Arg Trp Val Glu Leu Pro
35 40

<210> 25
<211> 38
<212> PRT
<213> *Caenorhabditis elegans*

<400> 25
Val Glu Tyr Glu Glu Ser Pro Ser Trp Leu Lys Leu Ile Tyr Tyr Glu
1 5 10 15
Glu Gly Thr Met Ile Gly Glu Lys Ala Asp Val Glu Gly His His Cys
20 25 30
Leu Ile Asp Gly Phe Thr
35

<210> 26
<211> 60
<212> PRT
<213> *Caenorhabditis elegans*

<400> 26
Asn Leu Ala Glu Thr Gly His Ser Lys Ile Met Arg Ala Ala His Lys
1 5 10 15
Val Ser Asn Pro Glu Ile Gly Tyr Cys Cys His Pro Thr Glu Tyr Asp
20 25 30
Tyr Ile Lys Leu Ile Tyr Val Asn Arg Asp Gly Arg Val Ser Ile Ala
35 40 45
Asn Val Asn Gly Met Ile Ala Lys Lys Cys Gly Cys
50 55 60

<210> 27
<211> 20
<212> PRT
<213> *Caenorhabditis elegans*

<400> 27
Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly
1 5 10 15
Asp Cys His Tyr
20

<210> 28
<211> 43
<212> PRT
<213> *Caenorhabditis elegans*

<400> 28
Val Cys Asn Ala Glu Ala Gln Ser Lys Gly Cys Cys Leu Tyr Asp Leu
1 5 10 15
Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp Trp Ile Val Ala Pro Pro

20 25 30
Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp Cys
35 40

<210> 29
<211> 70
<212> PRT
<213> *Caenorhabditis elegans*

<400> 29
Asp Cys His Tyr Asn Ala His His Phe Asn Leu Ala Glu Thr Gly His
1 5 10 15
Ser Lys Ile Met Arg Ala Ala His Lys Val Ser Asn Pro Glu Ile Gly
20 25 30
Tyr Cys Cys His Pro Thr Glu Tyr Asp Tyr Ile Lys Leu Ile Tyr Val
35 40 45
Asn Arg Asp Gly Arg Val Ser Ile Ala Asn Val Asn Gly Met Ile Ala
50 55 60
Lys Lys Cys Gly Cys Ser
65 70

<210> 30
<211> 35
<212> PRT
<213> *Caenorhabditis elegans*

<400> 30
Cys Cys Leu Tyr Asp Leu Glu Ile Glu Phe Glu Lys Ile Gly Trp Asp
1 5 10 15
Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala Tyr Met Cys Arg Gly Asp
20 25 30
Cys His Tyr
35

<210> 31
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Degenerate probe

<221> misc_feature
<222> (1)...(23)
<223> n = A,T,C or G

<400> 31
ggntggayt rnrtnrtngc ncc

23

<210> 32
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Degenerate probe

<221> misc_feature
<222> (1)...(18)
<223> n = A,T,C or G

<400> 32
tgytgynnnnc cnacngar

18

<210> 33
<211> 127
<212> PRT
<213> *Caenorhabditis elegans*

<400> 33
Lys Phe His Glu Trp Ala Ala Gln Ile Cys Asp Gly Met Ala Tyr Leu
1 5 10 15
Glu Ser Leu Lys Phe Cys His Arg Asp Leu Ala Ala Arg Asn Cys Met
20 25 30
Ile Asn Arg Asp Glu Thr Val Lys Ile Gly Asp Phe Gly Met Ala Arg
35 40 45
Asp Leu Phe Tyr His Asp Tyr Tyr Lys Pro Ser Gly Lys Arg Met Met
50 55 60
Pro Val Arg Trp Met Ser Pro Glu Ser Leu Lys Asp Gly Lys Phe Asp
65 70 75 80
Ser Lys Ser Asp Val Trp Ser Phe Gly Val Val Leu Tyr Glu Met Val
85 90 95
Thr Leu Gly Ala Gln Pro Tyr Ile Gly Leu Ser Asn Asp Glu Val Leu
100 105 110
Asn Tyr Ile Gly Met Ala Arg Lys Val Ile Lys Lys Pro Glu Cys
115 120 125

<210> 34
<211> 131
<212> PRT
<213> *Caenorhabditis elegans*

<400> 34
Asn Thr Thr Cys Gln Lys Ser Cys Ala Tyr Asp Arg Leu Leu Pro Thr
1 5 10 15
Lys Glu Ile Gly Pro Gly Cys Asp Ala Asn Gly Asp Arg Cys His Asp
20 25 30
Gln Cys Val Gly Gly Cys Glu Arg Val Asn Asp Ala Thr Ala Cys His
35 40 45
Ala Cys Lys Asn Val Tyr His Lys Gly Lys Cys Ile Glu Lys Cys Asp
50 55 60
Ala His Leu Tyr Leu Leu Gln Arg Arg Cys Val Thr Arg Glu Gln
65 70 75 80
Cys Leu Gln Leu Asn Pro Val Leu Ser Asn Lys Thr Val Pro Ile Lys
85 90 95
Ala Thr Ala Gly Leu Cys Ser Asp Lys Cys Pro Asp Gly Tyr Gln Ile
100 105 110
Asn Pro Asp Asp His Arg Glu Cys Arg Lys Cys Val Gly Lys Cys Glu
115 120 125
Ile Val Cys
130

<210> 35
<211> 103

<212> PRT

<213> *Caenorhabditis elegans*

<400> 35

Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys
1 5 10 15
Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr
20 25 30
Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu
35 40 45
Gln Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu
50 55 60
Trp Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His
65 70 75 80
Cys Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro
85 90 95
Tyr His Tyr Glu Ile Val Ile
100

<210> 36

<211> 79

<212> PRT

<213> *Caenorhabditis elegans*

<400> 36

Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val
1 5 10 15
Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr
20 25 30
Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val
35 40 45
Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys
50 55 60
Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
65 70 75

<210> 37

<211> 106

<212> PRT

<213> *Caenorhabditis elegans*

<400> 37

Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp Gly Asn Met Ser Tyr Ala
1 5 10 15
Glu Leu Ile Thr Thr Ala Ile Met Ala Ser Pro Glu Lys Arg Leu Thr
20 25 30
Leu Ala Gln Val Tyr Glu Trp Met Val Gln Asn Val Pro Tyr Phe Arg
35 40 45
Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly Trp Lys Asn Ser Ile Arg
50 55 60
His Asn Leu Ser Leu His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly
65 70 75 80
Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly
85 90 95
Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
100 105

<210> 38
 <211> 60
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 38
 Glu Ile Lys Leu Ser Asp Phe Lys His Gln Leu Phe Glu Leu Ile Ala
 1 5 10 15
 Pro Met Lys Trp Gly Thr Tyr Ser Val Lys Pro Gln Asp Tyr Val Phe
 20 25 30
 Arg Gln Leu Asn Asn Phe Gly Glu Ile Glu Val Ile Phe Asn Asp Asp
 35 40 45
 Gln Pro Leu Ser Lys Leu Glu Leu His Gly Thr Phe
 50 55 60

<210> 39
 <211> 2784
 <212> DNA
 <213> *Caenorhabditis elegans*

<400> 39
 atgaagctaa tagcaacttc tcttctagtt cccgacgagc acacaccgat gatgtcacca 60
 gtgaatacaa ctacaaagat tctacaacgg agtggattaa aatggaaat cccgccccat 120
 ttggatccag acagtcagga tcatgaccgg gaagatggtg tcaactaccc ggatccagat 180
 ttatggaca caaaaaacac aaatatgacc gactacgatt tggatgtgtt gaagcttgga 240
 aaaccagcag tagatgaagc acggaaaaag atcgaagttc ccgacgctag tgcgccccca 300
 aacaaaattt tagatattt gatgtattt agaacgttaa aagaaagtga actcatacaa 360
 ctgaatgcgt atcggacaaa acgaaatcga ttatcggttta acttggtcaa aaacaatatt 420
 gatcgagagt tcgaccaaaa agcttgcgag tccctgggtt aaaaatttggaa ggataagaag 480
 aatgatctcc agaacctgat tcatgtgggtt ctttcaaaag gtacaaaata taccgggtgc 540
 attacaattt caaggacact tcatggccgg ttacaggtcc acggaagaaa aggtttccct 600
 cacgtgtct atggcaact gtggagggtt aatgaaatga caaaaaacga aacgcgtcat 660
 gtggaccact gcaagcacgc atttggaaatg aaaaatgtaca tggatgcgtt gaatccctat 720
 cactacgaaa ttgtcattgg aactatgatt gttggcaga gggatcatga caatcgagat 780
 atgccgcgc cacatcaacg ctaccacact ccaggtcggc aggtccagt tgacgatatg 840
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 cagcctatgc ctcaacaatt gccttcagg ggcgcaacgt ttgcccattcc tctcccacat 960
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 catcaggat atgaaatgaa tgggcccagt tgctcttcgt aaaaacaacaa tccattccac 1140
 caaaaatcacc attataatga tattagccat ccaaataactt attcctacga ctgtggccg 1200
 aacttgcgtt ggtttccaaat tccttaccc gattttaccat atcctttcaa tcagcaacca 1260
 caccagccgc cacaactatc aaaaaaccat acgtcccaac aaggcagtca tcaaccagg 1320
 caccaggc acgttccgaa tcatggacca atttcaagac cagtgttaca accatcaaca 1380
 gtcacccgtt acgtgttccg tcgggtactgt agacagacat ttggaaatcg attttttgaa 1440
 ggagaaatgtt aacaatccgg cgcaataattt cgtcttagta acaaattcat tgaagaattt 1500
 gattcggcga tttgtgggtt gacagttgtt cgaccggcga tgacagacgg tgaggtttg 1560
 gagaacatca tgccggaga tgcaccatcat gacatgtt gcaagttcat tttgaggctc 1620
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 tgggaaacaa ttgtgtacta tgagaaaaat ttgcaattt gcgagaaaaaa atgttcgaga 1740
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 gagccaaatc caatttagaga accagtggcg tttaaagttc gtaaagcaat agtggatgga 1860
 attcgctttt cctacaaaaaa agacgggagt gttggctt aaaaaccgcat gaagtacccg 1920
 gtatgttca cttctgggtt tctcgacgag caatcaggag gcctaaagaa ggataaagtg 1980
 cacaatgtt acggatgtgc gtctatcaa acgtttggct tcaacgttcc caaacaatc 2040
 atcagagacg cgcttcttc caagcaaattt gcaacaatgtt acttgcaggaaaatttact 2100
 ccgatgaatt atatctacga gaagaagact caggaagagc tgcgaaggaa agcaacacgc 2160
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gaagcatacc	cagaacgccc	gtcaattcat	gattgtccag	tttggattga	gttgaaaatc	2280
aacattgcct	acgatttcat	ggattcaatc	tgccagtaca	taacccaactg	cttcgagccg	2340
ctaggaatgg	aagattttgc	aaaattggga	atcaacgtca	gtgatgacta	aatgataact	2400
ttttcactc	accctactag	atactgatt	agtcttattc	caaatcatcc	aacgatatca	2460
aacttttcc	tttgaacttt	gcatactatg	ttatcacaag	ttccaagcag	tttcaataca	2520
aacataggat	atgttaacaa	cttttgataa	gaatcaagtt	accaactgtt	cattgtgagc	2580
tttgagctgt	atagaaggac	aatgtatccc	atacctcaat	ctttaatagt	catcagtcac	2640
tggtcccgca	ccaattttt	cgattcgcata	atgtcatata	ttgcaccgtg	gccctttta	2700
ttgtaactt	taatatattt	tcttcccaac	ttgtgaatat	gattgatgaa	ccaccattt	2760
gagtaataaa	tgtatTTTT	gtgg				2784

<210> 40

<211> 796

<212> PRT

<213> *Caenorhabditis elegans*

<400> 40

Met	Lys	Leu	Ile	Ala	Thr	Ser	Leu	Leu	Val	Pro	Asp	Glu	His	Thr	Pro
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Met	Met	Ser	Pro	Val	Asn	Thr	Thr	Thr	Lys	Ile	Leu	Gln	Arg	Ser	Gly
									20					25	30
Ile	Lys	Met	Glu	Ile	Pro	Pro	Tyr	Leu	Asp	Pro	Asp	Ser	Gln	Asp	Asp
								35					40	45	
Asp	Pro	Glu	Asp	Gly	Val	Asn	Tyr	Pro	Asp	Pro	Asp	Leu	Phe	Asp	Thr
								50					55	60	
Lys	Asn	Thr	Asn	Met	Thr	Glu	Tyr	Asp	Leu	Asp	Val	Val	Lys	Leu	Gly
						65		70			75		80		
Lys	Pro	Ala	Val	Asp	Glu	Ala	Arg	Lys	Lys	Ile	Glu	Val	Pro	Asp	Ala
								85			90		95		
Ser	Ala	Pro	Pro	Asn	Lys	Ile	Val	Glu	Tyr	Leu	Met	Tyr	Tyr	Arg	Thr
								100			105		110		
Leu	Lys	Glu	Ser	Glu	Leu	Ile	Gln	Leu	Asn	Ala	Tyr	Arg	Thr	Lys	Arg
								115			120		125		
Asn	Arg	Leu	Ser	Leu	Asn	Leu	Val	Lys	Asn	Asn	Ile	Asp	Arg	Glu	Phe
								130			135		140		
Asp	Gln	Lys	Ala	Cys	Glu	Ser	Leu	Val	Lys	Lys	Leu	Lys	Asp	Lys	Lys
								145			150		155	160	
Asn	Asp	Leu	Gln	Asn	Leu	Ile	Asp	Val	Val	Leu	Ser	Lys	Gly	Thr	Lys
								165			170		175		
Tyr	Thr	Gly	Cys	Ile	Thr	Ile	Pro	Arg	Thr	Leu	Asp	Gly	Arg	Leu	Gln
								180			185		190		
Val	His	Gly	Arg	Lys	Gly	Phe	Pro	His	Val	Val	Tyr	Gly	Lys	Leu	Trp
								195			200		205		
Arg	Phe	Asn	Glu	Met	Thr	Lys	Asn	Glu	Thr	Arg	His	Val	Asp	His	Cys
								210			215		220		
Lys	His	Ala	Phe	Glu	Met	Lys	Ser	Asp	Met	Val	Cys	Val	Asn	Pro	Tyr
								225			230		235	240	
His	Tyr	Glu	Ile	Val	Ile	Gly	Thr	Met	Ile	Val	Gly	Gln	Arg	Asp	His
								245			250		255		
Asp	Asn	Arg	Asp	Met	Pro	Pro	Pro	His	Gln	Arg	Tyr	His	Thr	Pro	Gly
								260			265		270		
Arg	Gln	Asp	Pro	Val	Asp	Asp	Met	Ser	Arg	Phe	Ile	Pro	Pro	Ala	Ser
								275			280		285		
Ile	Arg	Pro	Pro	Pro	Met	Asn	Met	His	Thr	Arg	Pro	Gln	Pro	Met	Pro
								290			295		300		
Gln	Gln	Leu	Pro	Ser	Val	Gly	Ala	Thr	Phe	Ala	His	Pro	Leu	Pro	His
								305			310		315	320	
Gln	Ala	Pro	His	Asn	Pro	Gly	Val	Ser	His	Pro	Tyr	Ser	Ile	Ala	Pro
								325			330		335		

Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln
 340 345 350
 Met Pro Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly
 355 360 365
 Pro Ser Cys Ser Ser Glu Asn Asn Asn Pro Phe His Gln Asn His His
 370 375 380
 Tyr Asn Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro
 385 390 395 400
 Asn Leu Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe
 405 410 415
 Asn Gln Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser
 420 425 430
 Gln Gln Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp
 435 440 445
 Pro Pro Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp
 450 455 460
 Val Phe Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu
 465 470 475 480
 Gly Glu Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe
 485 490 495
 Ile Glu Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro
 500 505 510
 Arg Met Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala
 515 520 525
 Pro Tyr His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser
 530 535 540
 Val Thr Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys
 545 550 555 560
 Trp Gly Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys
 565 570 575
 Lys Cys Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser
 580 585 590
 Glu Asn Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro
 595 600 605
 Val Ala Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser
 610 615 620
 Tyr Lys Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro
 625 630 635 640
 Val Phe Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys
 645 650 655
 Lys Asp Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe
 660 665 670
 Gly Phe Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys
 675 680 685
 Gln Met Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr
 690 695 700
 Ile Tyr Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg
 705 710 715 720
 Thr Thr Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys
 725 730 735
 Lys Gly Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys
 740 745 750
 Pro Val Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp
 755 760 765
 Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu
 770 775 780
 Asp Phe Ala Lys Leu Gly Ile Asn Val Ser Asp Asp
 785 790 795

<210> 41
<211> 858
<212> PRT
<213> *Caenorhabditis elegans*

<400> 41
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Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu
20 25 30
Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu
35 40 45
Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu
50 55 60
Asn Val Gly Asn Met Ala Asn Val Pro Asp Glu His Thr Pro Met Met
65 70 75 80
Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly Ile Lys
85 90 95
Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp Asp Pro
100 105 110
Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr Lys Asn
115 120 125
Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly Lys Pro
130 135 140
Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala Ser Ala
145 150 155 160
Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr Leu Lys
165 170 175
Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg Asn Arg
180 185 190
Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe Asp Gln
195 200 205
Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys Asn Asp
210 215 220
Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys Tyr Thr
225 230 235 240
Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln Val His
245 250 255
Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp Arg Phe
260 265 270
Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys Lys His
275 280 285
Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr His Tyr
290 295 300
Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His Asp Asn
305 310 315 320
Arg Asp Met Pro Pro His Gln Arg Tyr His Thr Pro Gly Arg Gln
325 330 335
Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser Ile Arg
340 345 350
Pro Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro Gln Gln
355 360 365
Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His Gln Ala
370 375 380
Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro Gln Thr
385 390 395 400
His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln Met Pro
405 410 415
Gln Met Pro Pro Pro Leu His Gln Gly Tyr Gly Met Asn Gly Pro Ser

420	425	430
Cys Ser Ser Glu Asn Asn Asn Pro Phe His Gln Asn His His Tyr Asn		
435	440	445
Asp Ile Ser His Pro Asn His Tyr Ser Tyr Asp Cys Gly Pro Asn Leu		
450	455	460
Tyr Gly Phe Pro Thr Pro Tyr Pro Asp Phe His His Pro Phe Asn Gln		
465	470	475
Gln Pro His Gln Pro Pro Gln Leu Ser Gln Asn His Thr Ser Gln Gln		
485	490	495
Gly Ser His Gln Pro Gly His Gln Gly Gln Val Pro Asn Asp Pro Pro		
500	505	510
Ile Ser Arg Pro Val Leu Gln Pro Ser Thr Val Thr Leu Asp Val Phe		
515	520	525
Arg Arg Tyr Cys Arg Gln Thr Phe Gly Asn Arg Phe Phe Glu Gly Glu		
530	535	540
Ser Glu Gln Ser Gly Ala Ile Ile Arg Ser Ser Asn Lys Phe Ile Glu		
545	550	555
Glu Phe Asp Ser Pro Ile Cys Gly Val Thr Val Val Arg Pro Arg Met		
565	570	575
Thr Asp Gly Glu Val Leu Glu Asn Ile Met Pro Glu Asp Ala Pro Tyr		
580	585	590
His Asp Ile Cys Lys Phe Ile Leu Arg Leu Thr Ser Glu Ser Val Thr		
595	600	605
Phe Ser Gly Glu Gly Pro Glu Val Ser Asp Leu Asn Glu Lys Trp Gly		
610	615	620
Thr Ile Val Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys		
625	630	635
Ser Arg Gly Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn		
645	650	655
Arg Tyr Ser Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala		
660	665	670
Phe Lys Val Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys		
675	680	685
Lys Asp Gly Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe		
690	695	700
Val Thr Ser Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp		
705	710	715
Lys Val His Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe		
725	730	735
Asn Val Ser Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met		
740	745	750
Ala Thr Met Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr		
755	760	765
Glu Lys Lys Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr		
770	775	780
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly		
785	790	795
Phe Gly Glu Ala Tyr Pro Glu Arg Pro Ser Ile His Asp Cys Pro Val		
805	810	815
Trp Ile Glu Leu Lys Ile Asn Ile Ala Tyr Asp Phe Met Asp Ser Ile		
820	825	830
Cys Gln Tyr Ile Thr Asn Cys Phe Glu Pro Leu Gly Met Glu Asp Phe		
835	840	845
Ala Lys Leu Gly Ile Asn Val Ser Asp Asp		
850	855	

<210> 42

<211> 892

<212> PRT

<213> *Caenorhabditis elegans*

<400> 42

Met Gly Asp His His Asn Leu Thr Gly Leu Pro Gly Thr Ser Ile Pro
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Pro Gln Phe Asn Tyr Ser Gln Pro Gly Thr Ser Thr Gly Gly Pro Leu
20 25 30
Tyr Gly Gly Lys Pro Ser His Gly Leu Glu Asp Ile Pro Asp Val Glu
35 40 45
Glu Tyr Glu Arg Asn Leu Leu Gly Ala Gly Ala Gly Phe Asn Leu Leu
50 55 60
Asn Val Gly Asn Met Ala Asn Glu Phe Lys Pro Ile Ile Thr Leu Asp
65 70 75 80
Thr Lys Pro Pro Arg Asp Ala Asn Lys Ser Leu Ala Phe Asn Gly Gly
85 90 95
Leu Lys Leu Ile Thr Pro Lys Thr Glu Val Pro Asp Glu His Thr Pro
100 105 110
Met Met Ser Pro Val Asn Thr Thr Lys Ile Leu Gln Arg Ser Gly
115 120 125
Ile Lys Met Glu Ile Pro Pro Tyr Leu Asp Pro Asp Ser Gln Asp Asp
130 135 140
Asp Pro Glu Asp Gly Val Asn Tyr Pro Asp Pro Asp Leu Phe Asp Thr
145 150 155 160
Lys Asn Thr Asn Met Thr Glu Tyr Asp Leu Asp Val Leu Lys Leu Gly
165 170 175
Lys Pro Ala Val Asp Glu Ala Arg Lys Lys Ile Glu Val Pro Asp Ala
180 185 190
Ser Ala Pro Pro Asn Lys Ile Val Glu Tyr Leu Met Tyr Tyr Arg Thr
195 200 205
Leu Lys Glu Ser Glu Leu Ile Gln Leu Asn Ala Tyr Arg Thr Lys Arg
210 215 220
Asn Arg Leu Ser Leu Asn Leu Val Lys Asn Asn Ile Asp Arg Glu Phe
225 230 235 240
Asp Gln Lys Ala Cys Glu Ser Leu Val Lys Lys Leu Lys Asp Lys Lys
245 250 255
Asn Asp Leu Gln Asn Leu Ile Asp Val Val Leu Ser Lys Gly Thr Lys
260 265 270
Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr Leu Asp Gly Arg Leu Gln
275 280 285
Val His Gly Arg Lys Gly Phe Pro His Val Val Tyr Gly Lys Leu Trp
290 295 300
Arg Phe Asn Glu Met Thr Lys Asn Glu Thr Arg His Val Asp His Cys
305 310 315 320
Lys His Ala Phe Glu Met Lys Ser Asp Met Val Cys Val Asn Pro Tyr
325 330 335
His Tyr Glu Ile Val Ile Gly Thr Met Ile Val Gly Gln Arg Asp His
340 345 350
Asp Asn Arg Asp Met Pro Pro His Gln Arg Tyr His Thr Pro Gly
355 360 365
Arg Gln Asp Pro Val Asp Asp Met Ser Arg Phe Ile Pro Pro Ala Ser
370 375 380
Ile Arg Pro Pro Met Asn Met His Thr Arg Pro Gln Pro Met Pro
385 390 395 400
Gln Gln Leu Pro Ser Val Gly Ala Thr Phe Ala His Pro Leu Pro His
405 410 415
Gln Ala Pro His Asn Pro Gly Val Ser His Pro Tyr Ser Ile Ala Pro
420 425 430
Gln Thr His Tyr Pro Leu Asn Met Asn Pro Ile Pro Gln Met Pro Gln

435	440	445
Met Pro Gln Met Pro Pro	Leu His Gln Gly	Tyr Gly Met Asn Gly
450	455	460
Pro Ser Cys Ser Ser	Glu Asn Asn Asn	Pro Phe His Gln Asn His His
465	470	475
Tyr Asn Asp Ile Ser His	Pro Asn His	Tyr Ser Tyr Asp Cys Gly Pro
485	490	495
Asn Leu Tyr Gly Phe Pro Thr Pro	Tyr Pro Asp Phe His His	Pro Phe
500	505	510
Asn Gln Gln Pro His Gln Pro	Pro Gln Leu Ser Gln Asn His Thr Ser	
515	520	525
Gln Gln Gly Ser His Gln Pro	Gly His Gln Gly Gln Val Pro Asn Asp	
530	535	540
Pro Pro Ile Ser Arg Pro	Val Leu Gln Pro Ser	Thr Val Thr Leu Asp
545	550	555
Val Phe Arg Arg Tyr Cys Arg	Gln Thr Phe Gly Asn Arg Phe	Phe Glu
565	570	575
Gly Glu Ser Glu Gln Ser Gly Ala	Ile Ile Arg Ser Ser Asn Lys Phe	
580	585	590
Ile Glu Glu Phe Asp Ser Pro	Ile Cys Gly Val Thr Val Val Arg Pro	
595	600	605
Arg Met Thr Asp Gly Glu Val	Leu Glu Asn Ile	Met Pro Glu Asp Ala
610	615	620
Pro Tyr His Asp Ile Cys Lys	Phe Ile Leu Arg Leu Thr Ser Glu Ser	
625	630	635
Val Thr Phe Ser Gly Glu Gly	Pro Glu Val Ser Asp Leu Asn Glu Lys	
645	650	655
Trp Gly Thr Ile Val Tyr Tyr	Glu Lys Asn Leu Gln Ile Gly Glu Lys	
660	665	670
Lys Cys Ser Arg Gly Asn Phe	His Val Asp Gly Gly Phe Ile Cys Ser	
675	680	685
Glu Asn Arg Tyr Ser Leu Gly	Leu Glu Pro Asn Pro Ile Arg Glu Pro	
690	695	700
Val Ala Phe Lys Val Arg	Lys Ala Ile Val Asp Gly Ile Arg Phe Ser	
705	710	715
Tyr Lys Lys Asp Gly Ser Val	Trp Leu Gln Asn Arg Met Lys Tyr Pro	
725	730	735
Val Phe Val Thr Ser Gly Tyr	Leu Asp Glu Gln Ser Gly Gly Leu Lys	
740	745	750
Lys Asp Lys Val His Lys Val	Tyr Gly Cys Ala Ser Ile Lys Thr Phe	
755	760	765
Gly Phe Asn Val Ser Lys	Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys	
770	775	780
Gln Met Ala Thr Met Tyr	Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr	
785	790	795
Ile Tyr Glu Lys Lys Thr	Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg	
805	810	815
Thr Thr Asp Ser Leu Ala Lys	Tyr Cys Cys Val Arg Val Ser Phe Cys	
820	825	830
Lys Gly Phe Gly Glu Ala Tyr	Pro Glu Arg Pro Ser Ile His Asp Cys	
835	840	845
Pro Val Trp Ile Glu Leu Lys	Ile Asn Ile Ala Tyr Asp Phe Met Asp	
850	855	860
Ser Ile Cys Gln Tyr Ile Thr Asn Cys Phe	Glu Pro Leu Gly Met Glu	
865	870	875
Asp Phe Ala Lys Leu Gly Ile Asn Val	Ser Asp Asp	880
885	890	

<210> 43
 <211> 3499
 <212> DNA
 <213> *Caenorhabditis elegans*

<400> 43

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tcatcccctc	ctcttactcc	ttcttctcg	tccgctacta	ctgtatttc	tggacatcta	180
cctgtataca	caccagtggc	cagtcatctg	ccattacaat	ttcatcaatt	gacacttctt	240
caacaacaac	cgccgtcctc	attcactccc	gattcttctt	catcctcaac	atcgctgtct	300
ttggctgaaa	ttcccgaaga	cgttatgatg	gagatgctgg	tagatcaggg	aactgatgca	360
tcgtcatccg	cctccacgtc	cacctcatct	gttgcagat	tcggagcga	cacgttcatg	420
aatacaccgg	atgatgtat	gatgaatgat	gatatggaac	cgattcctcg	tgatcggtgc	480
aatacgtggc	caatgcgtag	gccgcaactc	gaaccaccac	tcaactcgag	tcccattatt	540
catgaacaaa	ttcctgaaga	agatgctgac	ctatacggga	gcaatgagca	atgtggacag	600
ctcggcggag	catcttcaaa	cgggtcgaca	gcaatgcttc	atactccaga	tggaagcaat	660
tctcatcaga	catcgttct	tcggagttc	agaatgtccg	aatcgccaga	cgataccgta	720
tcggaaaaaa	agacaacgac	cagacggAAC	gcttggggaa	atatgtcata	tgctgaactt	780
atcaactacag	ccattatggc	tagtccagag	aaacggtaa	ctcttgacca	agtttacgaa	840
tggatggtcc	agaatgttcc	atacttcagg	gataagggag	attcgaacag	ttcagctgga	900
tggaagaact	cgatccgtca	caatctgtct	cttcatttctc	gtttcatgca	aattcagaat	960
gaaggagccg	gaaagagctc	gtgggggtt	attaatccag	atgcaaagcc	aggaatgaat	1020
ccacggcgta	cacgtgaacg	atccaatact	attgagacga	ctacaaaggc	tcaactcgaa	1080
aaatctcgcc	gcggagccaa	gaagaggata	aaggagagag	cattgtggg	ctcccttcac	1140
tcgacactta	atgaaaattc	gattgccga	tcgattcaaa	cgatttctca	cgatttgcata	1200
gatgatgatc	aatgcaagga	gcatttgata	aegttccatc	atctttccgt	ccccgaactc	1260
aatcgaacct	ctcgatttct	ggatcgctgt	ctcgtgttc	tccagctatt	ggaagtgata	1320
tctatgatga	tctagaattt	ccatcatggg	ttggcgaatc	ggttccagca	attccaagtg	1380
atattgtga	tagaactgat	caaatgcgt	tcgatgcaac	tactcatgt	tggtggagtt	1440
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cacgagttga	acagtgtccg	tggatcggt	gctcagaatc	cacttcttcg	aaatccaatt	1560
gtgccaagca	ctaacttcaa	gccaatgcca	ctaccgggtg	cctatggaaa	ctatcaaaat	1620
ggtggaaataa	ctccaaatcaa	ttggctatca	acatccaact	catctccact	gcctggaatt	1680
caatcggtg	gaattgttagc	tgcacagcat	actgtcgctt	cttcatttcggc	tcttccaatt	1740
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ctcttcattt	tgttccctt	gggtttgttc	gaaagagaga	tagcaaagca	gcgaggagtg	1920
aggttaagcag	caataaaaaat	tttggatttt	tttttggttt	ttccagaaat	aatcgatttt	1980
ctggaaaatt	tcaaaaaaaaaa	atcggaattt	ttagttaaatt	atttgatgag	aaaaaaaaaaat	2040
tagaaaacat	aaggaaaaat	gaaaagcggtt	ttttttttc	gaaaattttt	gaattctcct	2100
acatttccaa	taagggcctt	agaactgcaa	acaaacaaaa	attgaaattt	tcgaatcaa	2160
aagttcccg	ataaaaagtag	ttcgaatatt	aaaaagcatt	taatttcctc	tttaaaaaaaaa	2220
ttgaataata	gccgaaattt	gcagattttt	tttctgaaaa	tcgaaaaacc	aaaatttttt	2280
gattttttaa	atttttttt	tacttccag	atagtaaaat	cattagcact	gaaaattatt	2340
tgaaaaaaaaaa	cttcaaatac	aaattttgtt	ttcgaaaaaaa	aaaatttaaa	tatatatttt	2400
cagaaatctt	ccgtcttcat	ctttcaaat	ccctacctac	acacactcaa	cgatcatcac	2460
agccagacca	tcaatattct	tccaaatttt	gacgtcgta	atttttttc	agttttttca	2520
aaaactctat	tttcttatttt	ctgtcgttt	ttccccttc	tctcgtaaa	ttccaacaca	2580
ttcatcccg	tgacgtcggt	taataataat	ataaaatacc	tcttctctt	ttcttccct	2640
aatgcgaaat	atcgaaaaac	cgttgattat	tacctctttt	ttcttgcgtt	ttttttctt	2700
ctctctctcc	cgtcatcccg	gttcttcaact	ctttaaatgc	tacctctatc	ccatctttt	2760
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aattgaattt	ttcaaaaaat	ttgatttctt	gatttctttt	gtaattcttt	aattttcttc	2880
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acatctggcc	gattcgaatc	ctccgtatac	acacacacat	agtaatctac	ctccaaaatt	3000
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gtatattgccc	accacgtcga	ttttaaattt	aaaccatcgt	tttttcttct	tttctacttt	3120
tttctcgaaa	aatttaacaa	cacacaaaaaa	aatccttcaa	aaaatctcag	ttttaaatgg	3180

<210> 45
<211> 510
<212> PRT
<213> *Caenorhabditis elegans*

<400> 45
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Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu Pro Ile Pro
35 40 45
Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln Leu Glu Pro
50 55 60
Pro Leu Asn Ser Ser Pro Ile Ile His Glu Gln Ile Pro Glu Glu Asp
65 70 75 80
Ala Asp Leu Tyr Gly Ser Asn Glu Gln Cys Gly Gln Leu Gly Gly Ala
85 90 95
Ser Ser Asn Gly Ser Thr Ala Met Leu His Thr Pro Asp Gly Ser Asn
100 105 110
Ser His Gln Thr Ser Phe Pro Ser Asp Phe Arg Met Ser Glu Ser Pro
115 120 125
Asp Asp Thr Val Ser Gly Lys Lys Thr Thr Thr Arg Arg Asn Ala Trp
130 135 140
Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser
145 150 155 160
Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln
165 170 175
Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
180 185 190
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met
195 200 205
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn
210 215 220
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser
225 230 235 240
Asn Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg
245 250 255
Gly Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His
260 265 270
Ser Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser
275 280 285
His Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val
290 295 300
Pro Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly
305 310 315 320
Ser Ser Ser Arg Val Ser Pro Ala Ile Gly Ser Asp Ile Tyr Asp Asp
325 330 335
Leu Glu Phe Pro Ser Trp Val Gly Glu Ser Val Pro Ala Ile Pro Ser
340 345 350
Asp Ile Val Asp Arg Thr Asp Gln Met Arg Ile Asp Ala Thr Thr His
355 360 365
Ile Gly Gly Val Gln Ile Lys Gln Glu Ser Lys Pro Ile Lys Thr Glu
370 375 380
Pro Ile Ala Pro Pro Pro Ser Tyr His Glu Leu Asn Ser Val Arg Gly
385 390 395 400
Ser Cys Ala Gln Asn Pro Leu Leu Arg Asn Pro Ile Val Pro Ser Thr
405 410 415
Asn Phe Lys Pro Met Pro Leu Pro Gly Ala Tyr Gly Asn Tyr Gln Asn

420	425	430	
Gly Gly Ile Thr Pro Ile Asn Trp	Leu Ser Thr Ser Asn Ser Ser Pro		
435	440	445	
Leu Pro Gly Ile Gln Ser Cys	Gly Ile Val Ala Ala Gln His Thr Val		
450	455	460	
Ala Ser Ser Ser Ala Leu Pro Ile Asp	Leu Glu Asn Leu Thr Leu Pro		
465	470	475	480
Asp Gln Pro Leu Met Asp Thr Met Asp	Val Asp Ala Leu Ile Arg His		
485	490	495	
Glu Leu Ser Gln Ala Gly Gly Gln His	Ile His Phe Asp Leu		
500	505	510	

<210> 46

<211> 509

<212> PRT

<213> *Caenorhabditis elegans*

<400> 46

Met Gln Gln Tyr Ile Tyr Gln Glu Ser Ser Ala Thr Ile Pro His His		
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Gln Leu Pro His Met Gln Gln Leu Pro Gln Pro Leu Leu Asn Leu Asn		
35 40 45		
Met Thr Thr Leu Thr Ser Ser Gly Ser Ser Val Ala Ser Ser Ile Gly		
50 55 60		
Gly Gly Ala Gln Cys Ser Pro Cys Ala Ser Gly Ser Ser Thr Ala Ala		
65 70 75 80		
Thr Asn Ser Ser Gln Gln Gln Gln Thr Val Gly Gln Met Leu Ala Ala		
85 90 95		
Ser Val Pro Cys Ser Ser Ser Gly Met Thr Leu Gly Met Ser Leu Asn		
100 105 110		
Leu Ser Gln Gly Gly Gly Pro Met Pro Ala Lys Lys Lys Arg Cys Arg		
115 120 125		
Lys Lys Pro Thr Asp Gln Leu Ala Gln Lys Lys Pro Asn Pro Trp Gly		
130 135 140		
Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys Ala Leu Glu Ser Ala Pro		
145 150 155 160		
Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr Gln Trp Phe Ser Asp Asn		
165 170 175		
Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro Glu Glu Ala Ala Gly Trp		
180 185 190		
Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met Arg		
195 200 205		
Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn Pro		
210 215 220		
Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg Glu Arg Ser Asn		
225 230 235 240		
Thr Ile Glu Thr Thr Lys Ala Gln Leu Glu Lys Ser Arg Arg Gly		
245 250 255		
Ala Lys Lys Arg Ile Lys Glu Arg Ala Leu Met Gly Ser Leu His Ser		
260 265 270		
Thr Leu Asn Gly Asn Ser Ile Ala Gly Ser Ile Gln Thr Ile Ser His		
275 280 285		
Asp Leu Tyr Asp Asp Asp Ser Met Gln Gly Ala Phe Asp Asn Val Pro		
290 295 300		
Ser Ser Phe Arg Pro Arg Thr Gln Ser Asn Leu Ser Ile Pro Gly Ser		
305 310 315 320		

Ser	Ser	Arg	Val	Ser	Pro	Ala	Ile	Gly	Ser	Asp	Ile	Tyr	Asp	Asp	Leu
				325				330				335			
Glu	Phe	Pro	Ser	Trp	Val	Gly	Glu	Ser	Val	Pro	Ala	Ile	Pro	Ser	Asp
				340				345				350			
Ile	Val	Asp	Arg	Thr	Asp	Gln	Met	Arg	Ile	Asp	Ala	Thr	Thr	His	Ile
				355				360				365			
Gly	Gly	Val	Gln	Ile	Lys	Gln	Glu	Ser	Lys	Pro	Ile	Lys	Thr	Glu	Pro
				370				375			380				
Ile	Ala	Pro	Pro	Pro	Ser	Tyr	His	Glu	Leu	Asn	Ser	Val	Arg	Gly	Ser
				385				390			395			400	
Cys	Ala	Gln	Asn	Pro	Leu	Leu	Arg	Asn	Pro	Ile	Val	Pro	Ser	Thr	Asn
				405				410			415				
Phe	Lys	Pro	Met	Pro	Leu	Pro	Gly	Ala	Tyr	Gly	Asn	Tyr	Gln	Asn	Gly
				420				425			430				
Gly	Ile	Thr	Pro	Ile	Asn	Trp	Leu	Ser	Thr	Ser	Asn	Ser	Ser	Pro	Leu
				435				440			445				
Pro	Gly	Ile	Gln	Ser	Cys	Gly	Ile	Val	Ala	Ala	Gln	His	Thr	Val	Ala
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Gln	Pro	Leu	Met	Asp	Thr	Met	Asp	Val	Asp	Ala	Leu	Ile	Arg	His	Glu
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<211> 3504

<212> DNA

<213> *Caenorhabditis elegans*

<400> 47

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<211> 1167

<212> PRT

<213> *Caenorhabditis elegans*

<400> 48

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Thr	Met	Val	Glu	Gln	Trp	Gln	Met	Arg	Glu	Arg	Pro	Ser	Leu	Glu	Thr
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Glu	Asn	Gly	Lys	Gly	Ser	Leu	Leu	Leu	Glu	Asn	Glu	Gly	Val	Ala	Asp
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							115		120			125			
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Ile Asn Arg Asp Lys Glu Leu Met Ser Asp Ile Ser His Cys Leu Gly
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 Tyr Ser Leu Asp Lys Leu Glu Glu Ser Leu Asp Glu Glu Leu Arg Gln
 180 185 190
 Phe Arg Ala Ser Leu Trp Ala Arg Thr Lys Lys Thr Cys Leu Thr Arg
 195 200 205
 Gly Leu Glu Gly Thr Ser His Tyr Ala Phe Pro Glu Glu Gln Tyr Leu
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 Cys Val Gly Glu Ser Cys Pro Lys Asp Leu Glu Ser Lys Val Lys Ala
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 Ala Lys Leu Ser Tyr Gln Met Phe Trp Arg Lys Arg Lys Ala Glu Ile
 245 250 255
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 Asn Glu Thr Pro Lys Ser Leu Leu His Thr Phe Leu Tyr Glu Met Arg
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 Lys Leu Asp Val Tyr Asp Thr Asp Asp Pro Ala Asp Glu Gly Trp Phe
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 Pro Gly Phe Val Val Arg Arg Gln Ser Leu Val Leu Lys Asp Tyr Cys
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 Lys Leu Ala Leu Asp Val Leu Ser Val Ser Ile Asp Ser Thr Pro Lys
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 Gln Ser Lys Asn Ser Asp Met Val Met Thr Asp Phe Arg Pro Thr Ala
 385 390 395 400
 Ser Leu Lys Gln Val Ser Leu Trp Asp Leu Asp Ala Asn Leu Met Ile
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 Arg Pro Val Asn Ile Ser Gly Phe Asp Phe Pro Ala Asp Val Asp Met
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 Tyr Val Arg Ile Glu Phe Ser Val Tyr Val Gly Thr Leu Thr Leu Ala
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 485 490 495
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 Glu Leu Arg Gln Gly Gln Phe Leu Phe His Leu Trp Ala Pro Glu Pro
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 530 535 540
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 610 615 620
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gtaccagcac	cggaggcccg	ctttaggtg	gaaaaccttc	tcatggattt	gaagatattc	180
ctgatgtaga	ggaatatgag	aggaacctgc	tcggggctgg	agcaggtttt	aatctgtca	240
atgttagaaaa	tatggctaat	gaatttaac	caataatcac	attggacacg	aaaccacctc	300
gtgatgcca	caagtcatgt	gcattcaatg	gcgggttcaa	gtaatcact	ccgaaaactg	360
aagttcccga	cgagcacaca	ccgatgtat	caccagtcaa	tacaactaca	aagattctac	420
aacggagtgg	tataaaatg	gaaatccgc	catatttgg	tccagacagt	caggatgt	480
acccggaaga	tgggtgtcaac	tacccggatc	cagatttt	tgacacaaaa	aacacaaata	540
tgaccgagta	cgatttggat	gtgttgaagc	ttggaaaaacc	agcagtagat	gaagcaccgga	600
aaaagatcga	agttcccgac	gctagtgcgc	cgcacaaacaa	aatttgtaa	tatgtatgt	660

attatagaac	gttaaaagaa	agtgaactca	tacaactgaa	tgcgtatcg	acaaaacgaa	720
atcgattatc	gttgaacttg	gtcaaaaaca	atattgatcg	agagttcgac	caaaaagctt	780
gcgagtccct	ggtgaaaaaa	ttgaaggata	agaagaatga	tctccagaac	ctgattgatg	840
tggttcttc	aaaaggtaca	aatataccg	gttgcattac	aattccaagg	acacttgatg	900
gccggttaca	ggtccacgga	agaaaagggt	tccctcacgt	agtctatggc	aaactgtgga	960
ggttaatga	aatgacaaaaa	aacgaaacgc	gtcatgtgga	ccactgcaag	cacgcatttg	1020
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tgattgttgg	gcagagggat	catgacaatc	gagatatgcc	gccgccacat	caacgctacc	1140
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ttcgtccgccc	tccgatgaac	atgcacacaa	ggcctcagcc	tatgcctcaa	caattgcctt	1260
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caccaatttc	aagaccagtg	ttacaaccat	caacagtac	cttggacgtg	ttccgtcggt	1740
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taattcggtc	tagtaacaaa	ttcattgaag	aatttggattc	gccgatttgt	gtgtgacag	1860
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catatcatga	cattgcaag	ttcattttga	ggctcacatc	agaaagtgt	actttctcag	1980
gagagggggcc	agaagttgt	gatttgaacg	aaaaatgggg	aacaattgt	tactatgaga	2040
aaaatttgc	aattggcgag	aaaaaatgtt	cgagaggaaa	tttccacgt	gatggcggt	2100
tcatttgctc	tgagaatcgt	tacagtctcg	gacttggcc	aaatccaatt	agagaaccag	2160
tggcgtttaa	agttcgtaaa	gcaatagtgg	atggaattcg	ctttccctac	aaaaaagacg	2220
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acgagcaatc	aggaggccta	aagaaggata	aagtgcacaa	agtttacgga	tgtgcgtcta	2340
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aaatggcaac	aatgtacttg	caaggaaaat	tgactccgt	gaattatatc	tacgagaaga	2460
agactcagga	agagctgcga	agggaaagcaa	cacgcaccac	tgattcattg	gccaagtact	2520
gttgtgtccg	tgtctcggtc	tgcaaaaggat	ttggagaagc	ataccagaa	cgcccgtcaa	2580
ttcatgattt	tccagtttg	attgagttga	aaatcaacat	tgcctacgt	ttcatggatt	2640
caatctgcca	gtacataacc	aactgcttcg	agccgctagg	aatggaagat	tttcaaaat	2700
tgggaatcaa	cgtcagtgt	gactaaatga	taacttttt	cactcaccc	actagatact	2760
gatttagtct	tattccaaat	catccaaacga	tatcaaactt	tttccttga	actttgcata	2820
ctatgttac	acaagttcca	agcagttca	atacaaacat	aggatatgtt	aacaacttt	2880
gataagaatc	aagttaccaa	ctgttcattg	tgagcttga	gctgtataga	aggacaatgt	2940
atcccataacc	tcaatcttta	atagtcatca	gtcactggtc	ccgcaccaat	tttttcgatt	3000
cgcataatgtc	atataattgca	ccgtggccct	tttatttga	acttttaata	tattttcttc	3060
ccaacttgc	aatatgattt	atgaaccacc	attttgagta	ataaatgtat	tttttgcgg	3119

<210> 54

<211> 103

<212> PRT

<213> *Caenorhabditis elegans*

<400> 54

Lys	Lys	Thr	Thr	Thr	Arg	Arg	Asn	Ala	Trp	Gly	Asn	Met	Ser	Tyr	Ala
1				5					10				15		
Glu	Leu	Ile	Thr	Thr	Ala	Ile	Met	Ala	Ser	Pro	Glu	Lys	Arg	Leu	Thr
				20				25				30			
Leu	Ala	Gln	Val	Tyr	Glu	Trp	Met	Val	Gln	Asn	Val	Pro	Tyr	Phe	Arg
				35				40				45			
Asp	Lys	Gly	Asp	Ser	Asn	Ser	Ser	Ala	Gly	Trp	Lys	Asn	Ser	Ile	Arg
	50				55					60					
His	Asn	Leu	Ser	Leu	His	Ser	Arg	Phe	Met	Arg	Ile	Gln	Asn	Glu	Gly
	65				70				75				80		
Ala	Gly	Lys	Ser	Ser	Trp	Trp	Val	Ile	Asn	Pro	Asp	Ala	Lys	Pro	Gly

85	90	95
Met Asn Pro Arg Arg Thr Arg		
100		

<210> 55	.	.
<211> 41		
<212> PRT		
<213> <i>Caenorhabditis elegans</i>		

<400> 55			
Thr Phe Met Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu			
1	5	10	15
Pro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Gln			
20		25	30
Leu Glu Pro Pro Leu Asn Ser Ser Pro			
35		40	

<210> 56		
<211> 109		
<212> PRT		
<213> <i>Caenorhabditis elegans</i>		

<400> 56				
Asp Asp Thr Val Ser Gly Lys Lys Thr Thr Arg Arg Asn Ala Trp				
1	5	10	15	
Gly Asn Met Ser Tyr Ala Glu Leu Ile Thr Thr Ala Ile Met Ala Ser				
20		25	30	
Pro Glu Lys Arg Leu Thr Leu Ala Gln Val Tyr Glu Trp Met Val Gln				
35		40	45	
Asn Val Pro Tyr Phe Arg Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly				
50		55	60	
Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Arg Phe Met				
65		70	75	80
Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser Trp Trp Val Ile Asn				
85		90		95
Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg Thr Arg				
100		105		

<210> 57		
<211> 655		
<212> PRT		
<213> <i>Homo sapiens</i>		

<400> 57				
Met Ala Glu Ala Pro Gln Val Val Glu Ile Asp Pro Asp Phe Glu Pro				
1	5	10	15	
Leu Pro Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro Arg Pro Glu Phe				
20		25	30	
Ser Gln Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro Ser Gly Ser Ala				
35		40	45	
Ala Ala Asn Pro Asp Ala Ala Gly Leu Pro Ser Ala Ser Ala Ala				
50		55	60	
Ala Val Ser Ala Asp Phe Met Ser Asn Leu Ser Leu Leu Glu Glu Ser				
65		70	75	80
Glu Asp Phe Pro Gln Ala Pro Gly Ser Val Ala Ala Ala Val Ala Ala				
85		90	95	

Ala Ala Ala Ala Ala Ala Thr Gly Gly Leu Cys Gly Asp Phe Gln Gly
 100 105 110
 Pro Glu Ala Gly Cys Leu His Pro Ala Pro Pro Gln Pro Pro Pro Pro
 115 120 125
 Gly Pro Val Ser Gln His Pro Pro Val Pro Pro Ala Ala Ala Gly Pro
 130 135 140
 Leu Ala Gly Gln Pro Arg Lys Ser Ser Ser Arg Arg Asn Ala Trp
 145 150 155 160
 Gly Asn Leu Ser Tyr Ala Asp Leu Ile Thr Lys Ala Ile Glu Ser Ser
 165 170 175
 Ala Glu Lys Arg Leu Thr Leu Ser Gln Ile Tyr Glu Trp Met Val Lys
 180 185 190
 Ser Val Pro Tyr Phe Lys Asp Lys Gly Asp Ser Asn Ser Ser Ala Gly
 195 200 205
 Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu His Ser Lys Phe Ile
 210 215 220
 Arg Val Gln Asn Glu Gly Thr Gly Lys Ser Ser Trp Trp Met Leu Asn
 225 230 235 240
 Pro Glu Gly Gly Lys Ser Gly Lys Ser Pro Arg Arg Arg Ala Ala Ser
 245 250 255
 Met Asp Asn Asn Ser Lys Phe Ala Lys Ser Arg Ser Arg Ala Ala Lys
 260 265 270
 Lys Lys Ala Ser Leu Gln Ser Gly Gln Glu Gly Ala Gly Asp Ser Pro
 275 280 285
 Gly Ser Gln Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His Ser Asn
 290 295 300
 Asp Asp Phe Asp Asn Trp Ser Thr Phe Arg Pro Arg Thr Ser Ser Asn
 305 310 315 320
 Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glu Gln Asp
 325 330 335
 Asp Leu Gly Glu Gly Asp Val His Ser Met Val Tyr Pro Pro Ser Ala
 340 345 350
 Ala Lys Met Ala Ser Thr Leu Pro Ser Leu Ser Glu Ile Ser Asn Pro
 355 360 365
 Glu Asn Met Glu Asn Leu Leu Asp Asn Leu Asn Leu Leu Ser Ser Pro
 370 375 380
 Thr Ser Leu Thr Val Ser Thr Gln Ser Ser Pro Gly Thr Met Met Gln
 385 390 395 400
 Gln Thr Pro Cys Tyr Ser Phe Ala Pro Pro Asn Thr Ser Leu Asn Ser
 405 410 415
 Pro Ser Pro Asn Tyr Gln Lys Tyr Thr Tyr Gly Gln Ser Ser Met Ser
 420 425 430
 Pro Leu Pro Gln Met Pro Ile Gln Thr Leu Gln Asp Asn Lys Ser Ser
 435 440 445
 Tyr Gly Gly Met Ser Gln Tyr Asn Cys Ala Pro Gly Leu Leu Lys Glu
 450 455 460
 Leu Leu Thr Ser Asp Ser Pro Pro His Asn Asp Ile Met Thr Pro Val
 465 470 475 480
 Asp Pro Gly Val Ala Gln Pro Asn Ser Arg Val Leu Gly Gln Asn Val
 485 490 495
 Met Met Gly Pro Asn Ser Val Met Ser Thr Tyr Gly Ser Gln Ala Ser
 500 505 510
 His Asn Lys Met Met Asn Pro Ser Ser His Thr His Pro Gly His Ala
 515 520 525
 Gln Gln Thr Ser Ala Val Asn Gly Arg Pro Leu Pro His Thr Val Ser
 530 535 540
 Thr Met Pro His Thr Ser Gly Met Asn Arg Leu Thr Gln Val Lys Thr
 545 550 555 560
 Pro Val Gln Val Pro Leu Pro His Pro Met Gln Met Ser Ala Leu Gly

	565	570	575												
Gly	Tyr	Ser	Ser	Val	Ser	Ser	Cys	Asn	Gly	Tyr	Gly	Arg	Met	Gly	Leu
580							585						590		
Leu	His	Gln	Glu	Lys	Leu	Pro	Ser	Asp	Leu	Asp	Gly	Met	Phe	Ile	Glu
595							600						605		
Arg	Leu	Asp	Cys	Asp	Met	Glu	Ser	Ile	Ile	Arg	Asn	Asp	Leu	Met	Asp
610						615							620		
Gly	Asp	Thr	Leu	Asp	Phe	Asn	Phe	Asp	Asn	Val	Leu	Pro	Asn	Gln	Ser
625						630				635				640	
Phe	Pro	His	Ser	Val	Lys	Thr	Thr	Thr	His	Ser	Trp	Val	Ser	Gly	
						645				650				655	

<210> 58
 <211> 98
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 58
 Lys Pro Asn Pro Trp Gly Glu Glu Ser Tyr Ser Asp Ile Ile Ala Lys
 1 5 10 15
 Ala Leu Glu Ser Ala Pro Asp Gly Arg Leu Lys Leu Asn Glu Ile Tyr
 20 25 30
 Gln Trp Phe Ser Asp Asn Ile Pro Tyr Phe Gly Glu Arg Ser Ser Pro
 35 40 45
 Glu Glu Ala Ala Gly Trp Lys Asn Ser Ile Arg His Asn Leu Ser Leu
 50 55 60
 His Ser Arg Phe Met Arg Ile Gln Asn Glu Gly Ala Gly Lys Ser Ser
 65 70 75 80
 Trp Trp Val Ile Asn Pro Asp Ala Lys Pro Gly Met Asn Pro Arg Arg
 85 90 95
 Thr Arg

<210> 59
 <211> 7
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 59
 Trp Lys Asn Ser Ile Arg His
 1 5

<210> 60
 <211> 121
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 60
 Gln Val Leu Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly
 1 5 10 15
 Val Gly Val Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr
 20 25 30
 Ser Lys Asp His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu
 35 40 45
 Arg Phe Pro Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly
 50 55 60

Leu Leu Val Lys Asp Pro Thr Gln Arg Leu Gly Gly Gly Pro Glu Asp
65 70 75 80
Ala Leu Glu Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu
85 90 95
Ala Thr Tyr Arg Lys Glu Ile Glu Pro Pro Tyr Lys Pro Asn Val Gln
100 105 110
Ser Glu Thr Asp Thr Ser Tyr Phe Asp
115 120

<210> 61
<211> 66
<212> PRT
<213> *Caenorhabditis elegans*

<400> 61
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
1 5 10 15
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
20 25 30
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
35 40 45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
50 55 60
Leu Thr
65

<210> 62
<211> 45
<212> PRT
<213> *Caenorhabditis elegans*

<400> 62
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 63
<211> 57
<212> PRT
<213> *Caenorhabditis elegans*

<400> 63
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1 5 10 15
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
20 25 30
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
35 40 45
Ile Val Leu Ala Leu Gly Tyr Leu His
50 55

<210> 64

<211> 59
<212> PRT
<213> *Caenorhabditis elegans*

<400> 64
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
1 5 10 15
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
35 40 45
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
50 55

<210> 65
<211> 33
<212> PRT
<213> *Caenorhabditis elegans*

<400> 65
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
1 5 10 15
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
20 25 30
Glu

<210> 66
<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 66
Val Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn
1 5 10 15
Trp Arg Pro Arg Phe
20

<210> 67
<211> 26
<212> PRT
<213> *Caenorhabditis elegans*

<400> 67
Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu Ile Val Leu Ala
1 5 10 15
Leu Gly Tyr Leu His Ala Asn Ser Ile Val
20 25

<210> 68
<211> 39
<212> PRT
<213> *Caenorhabditis elegans*

<400> 68

Ile Arg Val Ser Phe Cys Lys Gly Phe Gly Glu Thr Tyr Ser Arg Leu
1 5 10 15
Lys Val Val Asn Leu Pro Cys Trp Ile Glu Ile Ile Leu His Glu Pro
20 25 30
Ala Asp Glu Tyr Asp Thr Val
35

<210> 69
<211> 45
<212> PRT
<213> *Caenorhabditis elegans*

<400> 69
Ser Arg Asn Ser Lys Ser Ser Gln Ile Arg Asn Thr Val Gly Ala Gly
1 5 10 15
Ile Gln Leu Ala Tyr Glu Asn Gly Glu Leu Trp Leu Thr Val Leu Thr
20 25 30
Asp Gln Ile Val Phe Val Gln Cys Pro Phe Leu Asn Gln
35 40 45

<210> 70
<211> 29
<212> PRT
<213> *Caenorhabditis elegans*

<400> 70
Asn Glu Met Leu Asp Pro Glu Pro Lys Tyr Pro Lys Glu Glu Lys Pro
1 5 10 15
Trp Cys Thr Ile Phe Tyr Tyr Glu Leu Thr Val Arg Val
20 25

<210> 71
<211> 29
<212> PRT
<213> *Caenorhabditis elegans*

<400> 71
Gln Leu Gly Lys Ala Phe Glu Ala Lys Val Pro Thr Ile Thr Ile Asp
1 5 10 15
Gly Ala Thr Gly Ala Ser Asp Glu Cys Arg Met Ser Leu
20 25

<210> 72
<211> 105
<212> PRT
<213> *Caenorhabditis elegans*

<400> 72
Ser Pro Asp Asp Gly Leu Leu Asp Ser Ser Glu Glu Ser Arg Arg Arg
1 5 10 15
Gln Lys Thr Cys Arg Val Cys Gly Asp His Ala Thr Gly Tyr Asn Phe
20 25 30
Asn Val Ile Thr Cys Glu Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala
35 40 45
Leu Arg Pro Lys Glu Phe Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile

50	55	60
Asn Ser Val Ser Arg Arg	Phe Cys Gln Lys Cys	Arg Leu Arg Lys Cys
65	70	75
Phe Thr Val Gly Met Lys Lys Glu Trp	Ile Leu Asn Glu Glu Gln Leu	80
85	90	95
Arg Arg Arg Lys Asn Ser Arg Leu Asn		
100	105	

<210> 73
<211> 89
<212> PRT
<213> *Caenorhabditis elegans*

<400> 73

Leu Asp Ser Ser Glu Glu Ser Arg Arg	Gln Lys Thr Cys Arg Val	
1 5	10 15	
Cys Gly Asp His Ala Thr Gly Tyr Asn Phe Asn Val Ile	Thr Cys Glu	
20 25	30	
Ser Cys Lys Ala Phe Phe Arg Arg Asn Ala Leu Arg Pro	Lys Glu Phe	
35 40	45	
Lys Cys Pro Tyr Ser Glu Asp Cys Glu Ile Asn Ser Val	Ser Arg Arg	
50 55	60	
Phe Cys Gln Lys Cys Arg Leu Arg Lys Cys Phe Thr Val	Gly Met Lys	
65 70	75 80	
Lys Glu Trp Ile Leu Asn Glu Glu Gln		
85		

<210> 74
<211> 73
<212> PRT
<213> *Caenorhabditis elegans*

<400> 74

Asp Ile Met Asn Ile Met Asp Val Thr Met Arg Arg Phe Val	Lys Val	
1 5	10 15	
Ala Lys Gly Val Pro Ala Phe Arg Glu Val Ser Gln Glu	Gly Lys Phe	
20 25	30	
Ser Leu Leu Lys Gly Gly Met Ile Glu Met Leu Thr Val	Arg Gly Val	
35 40	45	
Thr Arg Tyr Asp Ala Ser Thr Asn Ser Phe Lys Thr Pro	Thr Ile Lys	
50 55	60	
Gly Gln Asn Val Ser Val Asn Val Asp		
65 70		

<210> 75
<211> 112
<212> PRT
<213> *Caenorhabditis elegans*

<400> 75

Ser Gly Ser Leu Val Asp Leu Met Ile Lys Asn Leu Thr Ala	Tyr Thr	
1 5	10 15	
Gln Gly Leu Asn Glu Thr Val Lys Asn Arg Thr Ala Glu	Leu Glu Lys	
20 25	30	
Glu Gln Glu Lys Gly Asp Gln Leu Leu Met Glu Leu Leu	Pro Lys Ser	
35 40	45	

Val Ala Asn Asp Leu Lys Asn Gly Ile Ala Val Asp Pro Lys Val Tyr
50 55 60
Glu Asn Ala Thr Ile Leu Tyr Ser Asp Ile Val Gly Phe Thr Ser Leu
65 70 75 80
Cys Ser Gln Ser Gln Pro Met Glu Val Val Thr Leu Leu Ser Gly Met
85 90 95
Tyr Gln Arg Phe Asp Leu Ile Ile Ser Gln Gln Gly Gly Tyr Lys Val
100 105 110

<210> 76
<211> 107
<212> PRT
<213> *Caenorhabditis elegans*

<400> 76
Met Glu Thr Ile Gly Asp Ala Tyr Cys Val Ala Ala Gly Leu Pro Val
1 5 10 15
Val Met Glu Lys Asp His Val Lys Ser Ile Cys Met Ile Ala Leu Leu
20 25 30
Gln Arg Asp Cys Leu His His Phe Glu Ile Pro His Arg Pro Gly Thr
35 40 45
Phe Leu Asn Cys Arg Trp Gly Phe Asn Ser Gly Pro Val Phe Ala Gly
50 55 60
Val Ile Gly Gln Lys Ala Pro Arg Tyr Ala Cys Phe Gly Glu Ala Val
65 70 75 80
Ile Leu Ala Ser Lys Met Glu Ser Ser Gly Val Glu Asp Arg Ile Gln
85 90 95
Met Thr Leu Ala Ser Gln Gln Leu Leu Glu Glu
100 105

<210> 77
<211> 43
<212> PRT
<213> *Caenorhabditis elegans*

<400> 77
Asp Ile Leu Lys Gly Leu Glu Tyr Ile His Ala Ser Ala Ile Asp Phe
1 5 10 15
His Gly Asn Leu Thr Leu His Asn Cys Met Leu Asp Ser His Trp Ile
20 25 30
Val Lys Leu Ser Gly Phe Gly Val Asn Arg Leu
35 40

<210> 78
<211> 15
<212> PRT
<213> *Caenorhabditis elegans*

<400> 78
Asp Met Tyr Ser Phe Gly Val Ile Leu His Glu Ile Ile Leu Lys
1 5 10 15

<210> 79
<211> 67
<212> PRT

<213> *Caenorhabditis elegans*

<400> 79

Ala	Ile	Lys	Ile	Asn	Val	Asp	Asp	Pro	Ala	Ser	Thr	Glu	Asn	Leu	Asn
1				5					10					15	
Tyr	Leu	Met	Glu	Ala	Asn	Ile	Met	Lys	Asn	Phe	Lys	Thr	Asn	Phe	Ile
						20		25					30		
Val	Gln	Leu	Tyr	Gly	Val	Ile	Ser	Thr	Val	Gln	Pro	Ala	Met	Val	Val
						35		40				45			
Met	Glu	Met	Met	Asp	Leu	Gly	Asn	Leu	Arg	Asp	Tyr	Leu	Arg	Ser	Lys
					50		55				60				
Arg	Glu	Asp													
		65													

<210> 80

<211> 54

<212> PRT

<213> *Caenorhabditis elegans*

<400> 80

Val	Ile	Lys	Lys	Pro	Glu	Cys	Cys	Glu	Asn	Tyr	Trp	Tyr	Lys	Val	Met
1				5					10				15		
Lys	Met	Cys	Trp	Arg	Tyr	Ser	Pro	Arg	Asp	Arg	Pro	Thr	Phe	Leu	Gln
				20				25				30			
Leu	Val	His	Leu	Leu	Ala	Ala	Glu	Ala	Ser	Pro	Glu	Phe	Arg	Asp	Leu
				35				40				45			
Ser	Phe	Val	Leu	Thr	Asp										
		50													

<210> 81

<211> 69

<212> PRT

<213> *Caenorhabditis elegans*

<400> 81

Lys	Gln	Asp	Ser	Gly	Met	Ala	Ser	Glu	Leu	Lys	Asp	Ile	Phe	Ala	Asn
1					5				10			15			
Ile	His	Thr	Ile	Thr	Gly	Tyr	Leu	Leu	Val	Arg	Gln	Ser	Ser	Pro	Phe
					20			25			30				
Ile	Ser	Leu	Asn	Met	Phe	Arg	Asn	Leu	Arg	Arg	Ile	Glu	Ala	Lys	Ser
				35				40			45				
Leu	Phe	Arg	Asn	Leu	Tyr	Ala	Ile	Thr	Val	Phe	Glu	Asn	Pro	Asn	Leu
				50				55			60				
Lys	Lys	Leu	Phe	Asp											
		65													

<210> 82

<211> 52

<212> PRT

<213> *Caenorhabditis elegans*

<400> 82

Phe	Pro	His	Leu	Arg	Glu	Ile	Thr	Gly	Thr	Leu	Leu	Val	Phe	Glu	Thr
1					5				10			15			
Glu	Gly	Leu	Val	Asp	Leu	Arg	Lys	Ile	Phe	Pro	Asn	Leu	Arg	Val	Ile
				20				25			30				

Gly Gly Arg Ser Leu Ile Gln His Tyr Ala Leu Ile Ile Tyr Arg Asn
35 40 45
Pro Asp Leu Glu
50

<210> 83
<211> 46
<212> PRT
<213> *Caenorhabditis elegans*

<400> 83
Glu Ile Gly Leu Asp Lys Leu Ser Val Ile Arg Asn Gly Gly Val Arg
1 5 10 15
Ile Ile Asp Asn Arg Lys Leu Cys Tyr Thr Lys Thr Ile Asp Trp Lys
20 25 30
His Leu Ile Thr Ser Ser Ile Asn Asp Val Val Val Asp Asn
35 40 45

<210> 84
<211> 36
<212> PRT
<213> *Caenorhabditis elegans*

<400> 84
Tyr Asn Ala Asp Asp Trp Glu Leu Arg Gln Asp Asp Val Val Leu Gly
1 5 10 15
Gln Gln Cys Gly Glu Gly Ser Phe Gly Lys Val Tyr Leu Gly Thr Gly
20 25 30
Asn Asn Val Val
35

<210> 85
<211> 24
<212> PRT
<213> *Caenorhabditis elegans*

<400> 85
Asp Ser Leu Ala Lys Tyr Cys Cys Val Arg Val Ser Phe Cys Lys Gly
1 5 10 15
Phe Gly Glu Ala Tyr Pro Glu Arg
20

<210> 86
<211> 13
<212> PRT
<213> *Caenorhabditis elegans*

<400> 86
Gly Trp Asp Trp Ile Val Ala Pro Pro Arg Tyr Asn Ala
1 5 10

<210> 87
<211> 121
<212> PRT

<213> Homo sapiens

<400> 87

Glu	Val	Leu	Glu	Asp	Asn	Asp	Tyr	Gly	Arg	Ala	Val	Asp	Trp	Trp	Gly
1			5					10						15	
Leu	Gly	Val	Val	Met	Tyr	Glu	Met	Met	Cys	Gly	Arg	Leu	Pro	Phe	Tyr
				20				25						30	
Asn	Gln	Asp	His	Glu	Lys	Leu	Phe	Glu	Leu	Ile	Leu	Met	Glu	Glu	Ile
					35			40					45		
Arg	Phe	Pro	Arg	Thr	Leu	Gly	Pro	Glu	Ala	Lys	Ser	Leu	Leu	Ser	Gly
					50			55				60			
Leu	Leu	Lys	Lys	Asp	Pro	Thr	Gln	Arg	Leu	Gly	Gly	Gly	Ser	Glu	Asp
					65			70		75			80		
Ala	Lys	Glu	Ile	Met	Gln	His	Arg	Phe	Phe	Ala	Asn	Ile	Val	Trp	Gln
					85			90				95			
Asp	Val	Tyr	Glu	Lys	Lys	Leu	Ser	Pro	Pro	Phe	Lys	Pro	Gln	Val	Thr
					100			105				110			
Ser	Glu	Thr	Asp	Thr	Arg	Tyr	Phe	Asp							
					115			120							

<210> 88

<211> 121

<212> PRT

<213> Caenorhabditis elegans

<400> 88

Gln	Val	Leu	Asp	Asp	His	Asp	Tyr	Gly	Arg	Cys	Val	Asp	Trp	Trp	Gly
1			5					10					15		
Val	Gly	Val	Val	Met	Tyr	Glu	Met	Met	Cys	Gly	Arg	Leu	Pro	Phe	Tyr
				20				25					30		
Ser	Lys	Asp	His	Asn	Lys	Leu	Phe	Glu	Leu	Ile	Met	Ala	Gly	Asp	Leu
					35			40				45			
Arg	Phe	Pro	Ser	Lys	Leu	Ser	Gln	Glu	Ala	Arg	Thr	Leu	Leu	Thr	Gly
					50			55			60				
Leu	Leu	Val	Lys	Asp	Pro	Thr	Gln	Arg	Leu	Gly	Gly	Gly	Pro	Glu	Asp
					65			70		75			80		
Ala	Leu	Glu	Ile	Cys	Arg	Ala	Asp	Phe	Phe	Arg	Thr	Val	Asp	Trp	Glu
					85			90				95			
Ala	Thr	Tyr	Arg	Lys	Glu	Ile	Glu	Pro	Pro	Tyr	Lys	Pro	Asn	Val	Gln
					100			105				110			
Ser	Glu	Thr	Asp	Thr	Ser	Tyr	Phe	Asp							
					115			120							

<210> 89

<211> 66

<212> PRT

<213> Homo sapiens

<400> 89

Thr	Met	Asn	Glu	Phe	Glu	Tyr	Leu	Lys	Leu	Leu	Gly	Lys	Gly	Thr	Phe
1				5					10				15		
Gly	Lys	Val	Ile	Leu	Val	Lys	Glu	Lys	Ala	Thr	Gly	Arg	Tyr	Tyr	Ala
					20			25				30			
Met	Lys	Ile	Leu	Lys	Lys	Glu	Val	Ile	Val	Ala	Lys	Asp	Glu	Val	Ala
					35			40				45			
His	Thr	Leu	Thr	Glu	Asn	Arg	Val	Leu	Gln	Asn	Ser	Arg	His	Pro	Phe
					50			55			60				

Leu Thr
65

<210> 90
<211> 66
<212> PRT
<213> *Caenorhabditis elegans*
<400> 90
Thr Met Glu Asp Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe
1 5 10 15
Gly Lys Val Ile Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala
20 25 30
Ile Lys Ile Leu Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala
35 40 45
His Thr Leu Thr Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe
50 55 60
Leu Thr
65

<210> 91
<211> 45
<212> PRT
<213> *Homo sapiens*
<400> 91
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 92
<211> 45
<212> PRT
<213> *Caenorhabditis elegans*
<400> 92
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 93
<211> 57
<212> PRT
<213> *Homo sapiens*
<400> 93
Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys
1 5 10 15
Phe Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser

20	25	30
Arg Glu Arg Val Phe Ser Glu Asp Arg Ala Arg Phe Tyr Gly Ala Glu		
35	40	45
Ile Val Ser Ala Leu Asp Tyr Leu His		
50	55	

(
<210> 94
<211> 57
<212> PRT
<213> *Caenorhabditis elegans*

<400> 94
Tyr Phe Gln Glu Leu Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys
1 5 10 15
Phe Val Met Gln Phe Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg
20 25 30
Lys Cys Gly Thr Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu
35 40 45
Ile Val Leu Ala Leu Gly Tyr Leu His
50 55

<210> 95
<211> 59
<212> PRT
<213> *Homo sapiens*

<400> 95
Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys Thr Glu Arg Pro
1 5 10 15
Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg Glu Glu Trp Ala
35 40 45
Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys
50 55

<210> 96
<211> 59
<212> PRT
<213> *Caenorhabditis elegans*

<400> 96
Ser Thr Phe Ala Ile Phe Tyr Phe Gln Thr Met Leu Phe Glu Lys Pro
1 5 10 15
Arg Pro Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile
20 25 30
Glu Arg Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile
35 40 45
His Ala Ile Glu Ser Ile Ser Lys Lys Tyr Lys
50 55

<210> 97
<211> 33
<212> PRT
<213> *Homo sapiens*

<400> 97
Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys Phe
1 5 10 15
Val Met Glu Tyr Ala Asn Gly Gly Glu Leu Phe Phe His Leu Ser Arg
20 25 30
Glu

<210> 98
<211> 33
<212> PRT
<213> *Caenorhabditis elegans*

<400> 98
Leu Gln Glu Leu Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe
1 5 10 15
Val Met Glu Phe Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg
20 25 30
Glu

<210> 99
<211> 36
<212> PRT
<213> *Homo sapiens* or *Caenorhabditis elegans*

<400> 99
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Asp Phe
1 5 10 15
Gly Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu
20 25 30
Ala Pro Glu Val
35

<210> 100
<211> 37
<212> PRT
<213> *Homo sapiens* or *Caenorhabditis elegans*

<400> 100
Leu Lys Tyr Ser Phe Gln Leu Cys Phe Val Met Ala Asn Gly Gly Glu
1 5 10 15
Leu Phe His Phe Ser Glu Arg Ala Arg Phe Tyr Gly Ala Glu Ile Val
20 25 30
Ala Leu Tyr Leu His
35

<210> 101
<211> 29
<212> PRT
<213> *Homo sapiens* or *Caenorhabditis elegans*

<400> 101
Phe Gln Met Glu Pro Arg Pro Asn Phe Arg Cys Leu Gln Trp Thr Thr
1 5 10 15

Val Ile Glu Arg Thr Phe Glu Glu Arg Trp Ala Ile Lys
20 25

<210> 102

<211> 24

<212> PRT

<213> Homo sapiens or Caenorhabditis elegans

<400> 102

Leu Leu Lys Tyr Ser Phe Gln Thr Asp Arg Leu Cys Phe Val Met Glu
1 5 10 15
Ala Gly Gly Leu His Leu Arg Glu
20

<210> 103

<211> 366

<212> PRT

<213> Homo sapiens

<400> 103

Arg Gly Ala Ile Arg Ile Glu Lys Asn Ala Asp Leu Cys Tyr Leu Ser
1 5 10 15
Thr Val Asp Trp Ser Leu Ile Leu Asp Ala Val Ser Asn Asn Tyr Ile
20 25 30
Val Gly Asn Lys Pro Pro Lys Glu Cys Gly Asp Leu Cys Pro Gly Thr
35 40 45
Met Glu Glu Lys Pro Met Cys Glu Lys Thr Thr Ile Asn Asn Glu Tyr
50 55 60
Asn Tyr Arg Cys Trp Thr Thr Asn Arg Cys Gln Lys Met Cys Pro Ser
65 70 75 80
Thr Cys Gly Lys Arg Ala Cys Thr Glu Asn Asn Glu Cys Cys His Pro
85 90 95
Glu Cys Leu Gly Ser Cys Ser Ala Pro Asp Asn Asp Thr Ala Cys Val
100 105 110
Ala Cys Arg His Tyr Tyr Tyr Ala Gly Val Cys Val Pro Ala Cys Pro
115 120 125
Pro Asn Thr Tyr Arg Phe Glu Gly Trp Arg Cys Val Asp Arg Asp Phe
130 135 140
Cys Ala Asn Ile Leu Ser Ala Glu Ser Ser Asp Ser Glu Gly Phe Val
145 150 155 160
Ile His Asp Gly Glu Cys Met Gln Glu Cys Pro Ser Gly Phe Ile Arg
165 170 175
Asn Gly Ser Gln Ser Met Tyr Cys Ile Pro Cys Glu Gly Pro Cys Pro
180 185 190
Lys Val Cys Glu Glu Lys Lys Thr Lys Thr Ile Asp Ser Val Thr
195 200 205
Ser Ala Gln Met Leu Gln Gly Cys Thr Ile Phe Lys Gly Asn Leu Leu
210 215 220
Ile Asn Ile Arg Arg Gly Asn Asn Ile Ala Ser Glu Leu Glu Asn Phe
225 230 235 240
Met Gly Leu Ile Glu Val Val Thr Gly Tyr Val Lys Ile Arg His Ser
245 250 255
His Ala Leu Val Ser Leu Ser Phe Leu Lys Asn Leu Arg Leu Ile Leu
260 265 270
Gly Glu Glu Gln Leu Glu Gly Asn Tyr Ser Phe Tyr Val Leu Asp Asn
275 280 285
Gln Asn Leu Gln Gln Leu Trp Asp Trp Asp His Arg Asn Leu Thr Ile

290	295	300
Lys Ala Gly Lys Met Tyr	Phe Ala Phe Asn Pro	Lys Leu Cys Val Ser
305	310	315
Glu Ile Tyr Arg Met Glu	Glu Val Thr Gly	Thr Lys Gly Arg Gln Ser
325	330	335
Lys Gly Asp Ile Asn Thr Arg Asn	Asn Gly Glu Arg Ala	Ser Cys Glu
340	345	350
Ser Asp Val Leu His Phe Thr	Ser Thr Thr Ser	Lys Asn
355	360	365

<210> 104
 <211> 370
 <212> PRT
 <213> Homo sapiens

<400> 104		
Arg Gly Ser Val Arg Ile Glu Lys Asn Asn Glu Leu Cys Tyr Leu Ala		
1	5	10
Thr Ile Asp Trp Ser Arg Ile Leu Asp Ser Val Glu Asp Asn Tyr Ile		
20	25	30
Val Leu Asn Lys Asp Asp Asn Glu Glu Cys Gly Asp Ile Cys Pro Gly		
35	40	45
Thr Ala Lys Gly Lys Thr Asn Cys Pro Ala Thr Val Ile Asn Gly Gln		
50	55	60
Phe Val Glu Arg Cys Trp Thr His Ser His Cys Gln Lys Val Cys Pro		
65	70	75
80		
Thr Ile Cys Lys Ser His Gly Cys Thr Ala Glu Gly Leu Cys Cys His		
85	90	95
Ser Glu Cys Leu Gly Asn Cys Ser Gln Pro Asp Asp Pro Thr Lys Cys		
100	105	110
Val Ala Cys Arg Asn Phe Tyr Leu Asp Gly Arg Cys Val Glu Thr Cys		
115	120	125
Pro Pro Pro Tyr Tyr His Phe Gln Asp Trp Arg Cys Val Asn Phe Ser		
130	135	140
Phe Cys Gln Asp Leu His His Lys Cys Lys Asn Ser Arg Arg Gln Gly		
145	150	155
160		
Cys His Gln Tyr Val Ile His Asn Asn Lys Cys Ile Pro Glu Cys Pro		
165	170	175
Ser Gly Tyr Thr Met Asn Ser Ser Asn Leu Leu Cys Thr Pro Cys Leu		
180	185	190
Gly Pro Cys Pro Lys Val Cys His Leu Leu Glu Gly Glu Lys Thr Ile		
195	200	205
Asp Ser Val Thr Ser Ala Gln Glu Leu Arg Gly Cys Thr Val Ile Asn		
210	215	220
Gly Ser Leu Ile Ile Asn Ile Arg Gly Gly Asn Asn Leu Ala Ala Glu		
225	230	235
240		
Leu Glu Ala Asn Leu Gly Leu Ile Glu Glu Ile Ser Gly Tyr Leu Lys		
245	250	255
Ile Arg Arg Ser Tyr Ala Leu Val Ser Leu Ser Phe Phe Arg Lys Leu		
260	265	270
Arg Leu Ile Arg Gly Glu Thr Leu Glu Ile Gly Asn Tyr Ser Phe Tyr		
275	280	285
Ala Leu Asp Asn Gln Asn Leu Arg Gln Leu Trp Asp Trp Ser Lys His		
290	295	300
Asn Leu Thr Ile Thr Gln Gly Lys Leu Phe Phe His Tyr Asn Pro Lys		
305	310	315
320		
Leu Cys Leu Ser Glu Ile His Lys Met Glu Glu Val Ser Gly Thr Lys		
325	330	335

Gly	Arg	Gln	Glu	Arg	Asn	Asp	Ile	Ala	Leu	Lys	Thr	Asn	Gly	Asp	Gln
			340						345						350
Ala	Ser	Cys	Glu	Asn	Glu	Leu	Leu	Lys	Phe	Ser	Tyr	Ile	Arg	Thr	Ser
			355				360								365
Phe	Asp														
			370												

<210> 105
<211> 383
<212> PRT
<213> Drosophila melanogaster

<400> 105															
Arg	Gly	Gly	Val	Arg	Ile	Glu	Lys	Asn	His	Lys	Leu	Cys	Tyr	Asp	Arg
1				5					10						15
Thr	Ile	Asp	Trp	Leu	Glu	Ile	Leu	Ala	Glu	Asn	Glu	Ser	Gln	Leu	Val
				20				25							30
Val	Leu	Thr	Glu	Asn	Gly	Lys	Glu	Lys	Glu	Cys	Ser	Leu	Ser	Lys	Cys
				35			40					45			
Pro	Gly	Glu	Ile	Arg	Ile	Glu	Glu	Gly	His	Asp	Asn	Thr	Ala	Ile	Glu
				50			55					60			
Gly	Glu	Leu	Asn	Ala	Ser	Cys	Gln	Leu	His	Asn	Asn	Arg	Arg	Leu	Cys
65					70				75						80
Trp	Asn	Ser	Lys	Leu	Cys	Gln	Thr	Lys	Cys	Pro	Glu	Lys	Cys	Arg	Asn
					85			90						95	
Asn	Cys	Ile	Asp	Glu	His	Thr	Cys	Cys	Ser	Gln	Asp	Cys	Leu	Gly	Gly
				100				105					110		
Cys	Val	Ile	Asp	Lys	Asn	Gly	Asn	Glu	Ser	Cys	Ile	Ser	Cys	Arg	Asn
				115				120				125			
Val	Ser	Phe	Asn	Asn	Ile	Cys	Met	Asp	Ser	Cys	Pro	Lys	Gly	Tyr	Tyr
					130		135				140				
Gln	Phe	Asp	Ser	Arg	Cys	Val	Thr	Ala	Asn	Glu	Cys	Ile	Thr	Leu	Thr
145						150				155					160
Lys	Phe	Glu	Thr	Asn	Ser	Val	Tyr	Ser	Gly	Ile	Pro	Tyr	Asn	Gly	Gln
						165			170						175
Cys	Ile	Thr	His	Cys	Pro	Thr	Gly	Tyr	Gln	Lys	Ser	Glu	Asn	Lys	Arg
					180			185				190			
Met	Cys	Glu	Pro	Cys	Pro	Gly	Gly	Lys	Cys	Asp	Lys	Glu	Cys	Ser	Ser
					195		200				205				
Gly	Leu	Ile	Asp	Ser	Leu	Glu	Arg	Ala	Arg	Glu	Phe	His	Gly	Cys	Thr
210						215					220				
Ile	Ile	Thr	Gly	Thr	Glu	Pro	Leu	Thr	Ile	Ser	Ile	Lys	Arg	Glu	Ser
225						230				235					240
Gly	Ala	His	Val	Met	Asp	Glu	Leu	Lys	Tyr	Gly	Leu	Ala	Ala	Val	His
					245				250						255
Lys	Ile	Gln	Ser	Ser	Leu	Met	Val	His	Leu	Thr	Tyr	Gly	Leu	Lys	Ser
					260			265				270			
Leu	Lys	Phe	Phe	Gln	Ser	Leu	Thr	Glu	Ile	Ser	Gly	Asp	Pro	Pro	Met
					275			280				285			
Asp	Ala	Asp	Lys	Tyr	Ala	Leu	Tyr	Val	Leu	Asp	Asn	Arg	Asp	Leu	Asp
					290		295				300				
Glu	Leu	Trp	Gly	Pro	Asn	Gln	Thr	Val	Phe	Ile	Arg	Lys	Gly	Gly	Val
305						310				315					320
Phe	Phe	His	Phe	Asn	Pro	Lys	Leu	Cys	Val	Ser	Thr	Ile	Asn	Gln	Leu
						325				330					335
Leu	Pro	Met	Leu	Ala	Ser	Lys	Pro	Lys	Phe	Phe	Glu	Lys	Ser	Asp	Glu
					340			345				350			
Gly	Ala	Asp	Ser	Asn	Gly	Asn	Arg	Gly	Ser	Cys	Gly	Thr	Ala	Val	Leu

Asn	Val	Thr	Leu	Gln	Ser	Val	Gly	Ala	Asn	Ser	Ala	Ser	Leu	Asn
355						360					365			
370						375					380			

<210> 106
 <211> 381
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 106															
Asn	Gly	Gly	Val	Arg	Ile	Ile	Asp	Asn	Arg	Lys	Leu	Cys	Tyr	Thr	Lys
1			5					10					15		
Thr	Ile	Asp	Trp	Lys	His	Leu	Ile	Thr	Ser	Ser	Ile	Asn	Asp	Val	Val
	20						25						30		
Val	Asp	Asn	Ala	Ala	Glu	Tyr	Ala	Val	Thr	Glu	Thr	Gly	Leu	Met	Cys
	35				40						45				
Pro	Arg	Gly	Ala	Cys	Glu	Glu	Asp	Lys	Gly	Glu	Ser	Lys	Cys	His	Tyr
	50				55			60							
Leu	Glu	Glu	Lys	Asn	Gln	Glu	Gln	Gly	Val	Glu	Arg	Val	Gln	Ser	Cys
65					70			75					80		
Trp	Ser	Asn	Thr	Thr	Cys	Gln	Lys	Ser	Cys	Ala	Tyr	Asp	Arg	Leu	Leu
					85			90					95		
Pro	Thr	Lys	Glu	Ile	Gly	Pro	Gly	Cys	Asp	Ala	Asn	Gly	Asp	Arg	Cys
	100					105						110			
His	Asp	Gln	Cys	Val	Gly	Gly	Cys	Glu	Arg	Val	Asn	Asp	Ala	Thr	Ala
	115					120						125			
Cys	His	Ala	Cys	Lys	Asn	Val	Tyr	His	Lys	Gly	Lys	Cys	Ile	Glu	Lys
	130					135			140						
Cys	Asp	Ala	His	Leu	Tyr	Leu	Leu	Gln	Arg	Arg	Cys	Val	Thr	Arg	
145					150				155				160		
Glu	Gln	Cys	Leu	Gln	Leu	Asn	Pro	Val	Leu	Ser	Asn	Lys	Thr	Val	Pro
					165				170				175		
Ile	Lys	Ala	Thr	Ala	Gly	Leu	Cys	Ser	Asp	Lys	Cys	Pro	Asp	Gly	Tyr
	180					185						190			
Gln	Ile	Asn	Pro	Asp	Asp	His	Arg	Glu	Cys	Arg	Lys	Cys	Val	Gly	Lys
	195					200						205			
Cys	Glu	Ile	Val	Cys	Glu	Ile	Asn	His	Val	Ile	Asp	Thr	Phe	Pro	Lys
210					215					220					
Ala	Gln	Ala	Ile	Arg	Leu	Cys	Asn	Ile	Ile	Asp	Gly	Asn	Leu	Thr	Ile
225					230				235					240	
Glu	Ile	Arg	Gly	Lys	Gln	Asp	Ser	Gly	Met	Ala	Ser	Glu	Leu	Lys	Asp
	245					250						255			
Ile	Phe	Ala	Asn	Ile	His	Thr	Ile	Thr	Gly	Tyr	Leu	Leu	Val	Arg	Gln
	260					265						270			
Ser	Ser	Pro	Phe	Ile	Ser	Leu	Asn	Met	Phe	Arg	Asn	Leu	Arg	Arg	Ile
275						280						285			
Glu	Ala	Lys	Ser	Leu	Phe	Arg	Asn	Leu	Tyr	Ala	Ile	Thr	Val	Phe	Glu
290					295					300					
Asn	Pro	Asn	Leu	Lys	Lys	Leu	Phe	Asp	Ser	Thr	Thr	Asp	Leu	Thr	Leu
305					310					315				320	
Asp	Arg	Gly	Thr	Val	Ser	Ile	Ala	Asn	Asn	Lys	Met	Leu	Cys	Phe	Lys
	325					330						335			
Tyr	Ile	Lys	Gln	Leu	Met	Ser	Lys	Leu	Asn	Ile	Pro	Leu	Asp	Pro	Ile
	340					345						350			
Asp	Gln	Ser	Glu	Gly	Thr	Asn	Gly	Glu	Lys	Ala	Ile	Cys	Glu	Asp	Met
	355					360						365			
Ala	Ile	Asn	Val	Ser	Ile	Thr	Ala	Val	Asn	Ala	Asp	Ser			
370						375					380				

<210> 107
<211> 370
<212> PRT
<213> Homo sapiens

<400> 107
Ala Leu Pro Val Ala Val Leu Leu Ile Val Gly Gly Leu Val Ile Met
1 5 10 15
Leu Tyr Val Phe His Arg Lys Arg Asn Asn Ser Arg Leu Gly Asn Gly
20 25 30
Val Leu Tyr Ala Ser Val Asn Pro Glu Tyr Phe Ser Ala Ala Asp Val
35 40 45
Tyr Val Pro Asp Glu Trp Glu Val Ala Arg Glu Lys Ile Thr Met Ser
50 55 60
Arg Glu Leu Gly Gln Gly Ser Phe Gly Met Val Tyr Glu Gly Val Ala
65 70 75 80
Lys Gly Val Val Lys Asp Glu Pro Glu Thr Arg Val Ala Ile Lys Thr
85 90 95
Val Asn Glu Ala Ala Ser Met Arg Glu Arg Ile Glu Phe Leu Asn Glu
100 105 110
Ala Ser Val Met Lys Glu Phe Asn Cys His His Val Val Arg Leu Leu
115 120 125
Gly Val Val Ser Gln Gly Gln Pro Thr Leu Val Ile Met Glu Leu Met
130 135 140
Thr Arg Gly Asp Leu Lys Ser Tyr Leu Arg Ser Leu Arg Pro Glu Met
145 150 155 160
Glu Asn Asn Pro Val Leu Ala Pro Pro Ser Leu Ser Lys Met Ile Gln
165 170 175
Met Ala Gly Glu Ile Ala Asp Gly Met Ala Tyr Leu Asn Ala Asn Lys
180 185 190
Phe Val His Arg Asp Leu Ala Ala Arg Asn Cys Met Val Ala Glu Asp
195 200 205
Phe Thr Val Lys Ile Gly Asp Phe Gly Met Thr Arg Asp Ile Tyr Glu
210 215 220
Thr Asp Tyr Tyr Arg Lys Gly Gly Lys Gly Leu Leu Pro Val Arg Trp
225 230 235 240
Met Ser Pro Glu Ser Leu Lys Asp Gly Val Phe Thr Thr Tyr Ser Asp
245 250 255
Val Trp Ser Phe Gly Val Val Leu Trp Glu Ile Ala Thr Leu Ala Glu
260 265 270
Gln Pro Tyr Gln Gly Leu Ser Asn Glu Gln Val Leu Arg Phe Val Met
275 280 285
Glu Gly Gly Leu Leu Asp Lys Pro Asp Asn Cys Pro Asp Met Leu Phe
290 295 300
Glu Leu Met Arg Met Cys Trp Gln Tyr Asn Pro Lys Met Arg Pro Ser
305 310 315 320
Phe Leu Glu Ile Ile Ser Ser Ile Lys Glu Glu Met Glu Pro Gly Phe
325 330 335
Arg Glu Val Ser Phe Tyr Tyr Ser Glu Glu Asn Lys Leu Pro Glu Pro
340 345 350
Glu Glu Leu Asp Leu Glu Pro Glu Asn Met Glu Ser Val Pro Leu Asp
355 360 365
Pro Ser
370

<210> 108
<211> 374

<212> PRT
<213> Homo sapiens

<400> 108

Ile	Gly	Pro	Leu	Ile	Phe	Val	Phe	Leu	Phe	Ser	Val	Val	Ile	Gly	Ser
1					5				10				15		
Ile	Tyr	Leu	Phe	Leu	Arg	Lys	Arg	Gln	Pro	Asp	Gly	Pro	Leu	Gly	Pro
					20			25					30		
Leu	Tyr	Ala	Ser	Ser	Asn	Pro	Glu	Tyr	Leu	Ser	Ala	Ser	Asp	Val	Phe
					35			40				45			
Pro	Cys	Ser	Val	Tyr	Val	Pro	Asp	Glu	Trp	Glu	Val	Ser	Arg	Glu	Lys
					50			55			60				
Ile	Thr	Leu	Leu	Arg	Glu	Leu	Gly	Gln	Gly	Ser	Phe	Gly	Met	Val	Tyr
					65			70		75			80		
Glu	Gly	Asn	Ala	Arg	Asp	Ile	Ile	Lys	Gly	Glu	Ala	Glu	Thr	Arg	Val
					85				90			95			
Ala	Val	Lys	Thr	Val	Asn	Glu	Ser	Ala	Ser	Leu	Arg	Glu	Arg	Ile	Glu
					100			105			110				
Phe	Leu	Asn	Glu	Ala	Ser	Val	Met	Lys	Gly	Phe	Thr	Cys	His	His	Val
					115			120			125				
Val	Arg	Leu	Leu	Gly	Val	Val	Ser	Lys	Gly	Gln	Pro	Thr	Leu	Val	Val
					130			135			140				
Met	Glu	Leu	Met	Ala	His	Gly	Asp	Leu	Lys	Ser	Tyr	Leu	Arg	Ser	Leu
					145			150		155			160		
Arg	Pro	Glu	Ala	Glu	Asn	Asn	Pro	Gly	Arg	Pro	Pro	Pro	Thr	Leu	Gln
					165				170			175			
Glu	Met	Ile	Gln	Met	Ala	Ala	Glu	Ile	Ala	Asp	Gly	Met	Ala	Tyr	Leu
					180			185			190				
Asn	Ala	Lys	Lys	Phe	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Cys	Met
					195			200			205				
Val	Ala	His	Asp	Phe	Thr	Val	Lys	Ile	Gly	Asp	Phe	Gly	Met	Thr	Arg
					210			215			220				
Asp	Ile	Tyr	Glu	Thr	Asp	Tyr	Tyr	Arg	Lys	Gly	Gly	Lys	Gly	Leu	Leu
					225			230		235			240		
Pro	Val	Arg	Trp	Met	Ala	Pro	Glu	Ser	Leu	Lys	Asp	Gly	Val	Phe	Thr
					245				250			255			
Thr	Ser	Ser	Asp	Met	Trp	Ser	Phe	Gly	Val	Val	Leu	Trp	Glu	Ile	Thr
					260			265			270				
Ser	Leu	Ala	Glu	Gln	Pro	Tyr	Gln	Gly	Leu	Ser	Asn	Glu	Gln	Val	Leu
					275			280			285				
Lys	Phe	Val	Met	Asp	Gly	Gly	Tyr	Leu	Asp	Gln	Pro	Asp	Asn	Cys	Pro
					290			295			300				
Glu	Arg	Val	Thr	Asp	Leu	Met	Arg	Met	Cys	Trp	Gln	Phe	Asn	Pro	Lys
					305			310		315			320		
Met	Arg	Pro	Thr	Phe	Leu	Glu	Ile	Val	Asn	Leu	Leu	Lys	Asp	Asp	Leu
					325				330			335			
His	Pro	Ser	Phe	Pro	Glu	Val	Ser	Phe	Phe	His	Ser	Glu	Glu	Asn	Lys
					340			345			350				
Ala	Pro	Glu	Ser	Glu	Glu	Leu	Glu	Met	Glu	Phe	Glu	Asp	Met	Glu	Asn
					355			360			365				
Val	Pro	Leu	Asp	Arg	Ser										
					370										

<210> 109
<211> 384
<212> PRT
<213> Drosophila melanogaster

<400> 109

Gly	Ile	Gly	Leu	Ala	Phe	Leu	Ile	Val	Ser	Leu	Phe	Gly	Tyr	Val	Cys
1															15
Tyr	Leu	His	Lys	Arg	Lys	Val	Pro	Ser	Asn	Asp	Leu	His	Met	Asn	Thr
															30
Glu	Val	Asn	Pro	Phe	Tyr	Ala	Ser	Met	Gln	Tyr	Ile	Pro	Asp	Asp	Trp
															45
Glu	Val	Leu	Arg	Glu	Asn	Ile	Ile	Gln	Leu	Ala	Pro	Leu	Gly	Gln	Gly
															60
Ser	Phe	Gly	Met	Val	Tyr	Glu	Gly	Ile	Leu	Lys	Ser	Phe	Pro	Pro	Asn
															80
Gly	Val	Asp	Arg	Glu	Cys	Ala	Ile	Lys	Thr	Val	Asn	Glu	Asn	Ala	Thr
															95
Asp	Arg	Glu	Arg	Thr	Asn	Phe	Leu	Ser	Glu	Ala	Ser	Val	Met	Lys	Glu
															110
Phe	Asp	Thr	Tyr	His	Val	Val	Arg	Leu	Leu	Gly	Val	Cys	Ser	Arg	Gly
															125
Gln	Pro	Ala	Leu	Val	Val	Met	Glu	Leu	Met	Lys	Lys	Gly	Asp	Leu	Lys
															140
Ser	Tyr	Leu	Arg	Ala	His	Arg	Pro	Glu	Glu	Arg	Asp	Glu	Ala	Met	Met
															160
Thr	Tyr	Leu	Asn	Arg	Ile	Gly	Val	Thr	Gly	Asn	Val	Gln	Pro	Pro	Thr
															175
Tyr	Gly	Arg	Ile	Tyr	Gln	Met	Ala	Ile	Glu	Ile	Ala	Asp	Gly	Met	Ala
															190
Tyr	Leu	Ala	Ala	Lys	Lys	Phe	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn
															205
Cys	Met	Val	Ala	Asp	Asp	Leu	Thr	Val	Lys	Ile	Gly	Asp	Phe	Gly	Met
															220
Thr	Arg	Asp	Ile	Tyr	Glu	Thr	Asp	Tyr	Tyr	Arg	Lys	Gly	Thr	Lys	Gly
															240
Leu	Leu	Pro	Val	Arg	Trp	Met	Pro	Pro	Glu	Ser	Leu	Arg	Asp	Gly	Val
															255
Tyr	Ser	Ser	Ala	Ser	Asp	Val	Phe	Ser	Phe	Gly	Val	Val	Leu	Trp	Glu
															270
Met	Ala	Thr	Leu	Ala	Ala	Gln	Pro	Tyr	Gln	Gly	Leu	Ser	Asn	Glu	Gln
															285
Val	Leu	Arg	Tyr	Val	Ile	Asp	Gly	Gly	Val	Met	Glu	Arg	Pro	Glu	Asn
															300
Cys	Pro	Asp	Phe	Leu	His	Lys	Leu	Met	Gln	Arg	Cys	Trp	His	His	Arg
															320
Ser	Ser	Ala	Arg	Pro	Ser	Phe	Leu	Asp	Ile	Ile	Ala	Tyr	Leu	Glu	Pro
															335
Gln	Cys	Pro	Asn	Ser	Gln	Phe	Lys	Glu	Val	Ser	Phe	Tyr	His	Ser	Glu
															350
Ala	Gly	Leu	Gln	His	Arg	Glu	Lys	Glu	Arg	Lys	Glu	Arg	Asn	Gln	Leu
															365
Asp	Ala	Phe	Ala	Ala	Val	Pro	Leu	Asp	Gln	Asp	Leu	Gln	Asp	Arg	Glu
															380

<210> 110

<211> 380

<212> PRT

<213> *Caenorhabditis elegans*

<400> 110

Gly	Met	Leu	Leu	Val	Phe	Leu	Ile	Leu	Met	Ser	Ile	Ala	Gly	Cys	Ile
1															15

Ile	Tyr	Tyr	Tyr	Ile	Gln	Val	Arg	Tyr	Gly	Lys	Lys	Val	Lys	Ala	Leu
				20				25				30			
Ser	Asp	Phe	Met	Gln	Leu	Asn	Pro	Glu	Tyr	Cys	Val	Asp	Asn	Lys	Tyr
				35				40			45				
Asn	Ala	Asp	Asp	Trp	Glu	Leu	Arg	Gln	Asp	Asp	Val	Val	Leu	Gly	Gln
				50				55			60				
Gln	Cys	Gly	Glu	Gly	Ser	Phe	Gly	Lys	Val	Tyr	Leu	Gly	Thr	Gly	Asn
				65				70		75			80		
Asn	Val	Val	Ser	Leu	Met	Gly	Asp	Arg	Phe	Gly	Pro	Cys	Ala	Ile	Lys
				85				90			95				
Ile	Asn	Val	Asp	Asp	Pro	Ala	Ser	Thr	Glu	Asn	Leu	Asn	Tyr	Leu	Met
				100				105			110				
Glu	Ala	Asn	Ile	Met	Lys	Asn	Phe	Lys	Thr	Asn	Phe	Ile	Val	Gln	Leu
				115				120			125				
Tyr	Gly	Val	Ile	Ser	Thr	Val	Gln	Pro	Ala	Met	Val	Val	Met	Glu	Met
				130				135			140				
Met	Asp	Leu	Gly	Asn	Leu	Arg	Asp	Tyr	Leu	Arg	Ser	Lys	Arg	Glu	Asp
				145				150		155			160		
Glu	Val	Phe	Asn	Glu	Thr	Asp	Cys	Asn	Phe	Phe	Asp	Ile	Ile	Pro	Arg
				165				170			175				
Asp	Lys	Phe	His	Glu	Trp	Ala	Ala	Gln	Ile	Cys	Asp	Gly	Met	Ala	Tyr
				180				185			190				
Leu	Glu	Ser	Leu	Lys	Phe	Cys	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Cys
				195				200			205				
Met	Ile	Asn	Arg	Asp	Glu	Thr	Val	Lys	Ile	Gly	Asp	Phe	Gly	Met	Ala
				210				215			220				
Arg	Asp	Leu	Phe	Tyr	His	Asp	Tyr	Tyr	Lys	Pro	Ser	Gly	Lys	Arg	Met
				225				230		235			240		
Met	Pro	Val	Arg	Trp	Met	Ser	Pro	Glu	Ser	Leu	Lys	Asp	Gly	Lys	Phe
				245				250			255				
Asp	Ser	Lys	Ser	Asp	Val	Trp	Ser	Phe	Gly	Val	Val	Leu	Tyr	Glu	Met
				260				265			270				
Val	Thr	Leu	Gly	Ala	Gln	Pro	Tyr	Ile	Gly	Leu	Ser	Asn	Asp	Glu	Val
				275				280			285				
Leu	Asn	Tyr	Ile	Gly	Met	Ala	Arg	Lys	Val	Ile	Lys	Lys	Pro	Glu	Cys
				290				295			300				
Cys	Glu	Asn	Tyr	Trp	Tyr	Lys	Val	Met	Lys	Met	Cys	Trp	Arg	Tyr	Ser
				305				310		315			320		
Pro	Arg	Asp	Arg	Pro	Thr	Phe	Leu	Gln	Leu	Val	His	Leu	Leu	Ala	Ala
				325				330			335				
Glu	Ala	Ser	Pro	Glu	Phe	Arg	Asp	Leu	Ser	Phe	Val	Leu	Thr	Asp	Asn
				340				345			350				
Gln	Met	Ile	Leu	Asp	Asp	Ser	Glu	Ala	Leu	Asp	Leu	Asp	Asp	Ile	Asp
				355				360			365				
Asp	Thr	Asp	Met	Asn	Asp	Gln	Val	Val	Glu	Val	Ala				
				370				375			380				

<210> 111
 <211> 103
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 111
 Asn Ile Asp Arg Glu Phe Asp Gln Lys Ala Cys Glu Ser Leu Val Lys
 1 5 10 15
 Lys Leu Lys Asp Lys Lys Asn Asp Leu Gln Asn Leu Ile Asp Val Val
 20 25 30
 Leu Ser Lys Gly Thr Lys Tyr Thr Gly Cys Ile Thr Ile Pro Arg Thr

35	40	45	
Leu Asp Gly Arg Leu Gln Val His	Gly Arg Lys Gly	Phe Pro His Val	
50	55	60	
Val Tyr Gly Lys Leu Trp Arg Phe Asn Glu Met	Thr Lys Asn Glu	Thr	
65	70	75	80
Arg His Val Asp His Cys Lys His Ala Phe	Glu Met Lys Ser Asp	Met	
85		90	95
Val Cys Val Asn Pro Tyr His			
100			

<210> 112
 <211> 104
 <212> PRT
 <213> Homo sapiens

<400> 112		
Gly	Gly	Glu Ser Glu Thr Phe Ala Lys Arg Ala Ile Glu Ser Leu Val
1	5	10 15
Lys	Lys	Leu Lys Glu Lys Lys Asp Glu Leu Asp Ser Leu Ile Thr Ala
	20	25 30
Ile	Thr	Thr Asn Gly Ala His Pro Ser Lys Cys Val Thr Ile Gln Arg
	35	40 45
Thr	Leu	Asp Gly Arg Leu Gln Val Ala Gly Arg Lys Gly Phe Pro His
	50	55 60
Val	Ile	Tyr Ala Arg Leu Trp Arg Trp Pro Asp Leu His Lys Asn Glu
	65	70 75 80
Leu	Lys	His Val Lys Tyr Cys Gln Tyr Ala Phe Asp Leu Lys Cys Asp
	85	90 95
Ser	Val	Cys Val Asn Pro Tyr His
	100	

<210> 113
 <211> 205
 <212> PRT
 <213> Caenorhabditis elegans

<400> 113		
Ile	Val	Tyr Tyr Glu Lys Asn Leu Gln Ile Gly Glu Lys Lys Cys Ser
1	5	10 15
Arg	Gly	Asn Phe His Val Asp Gly Gly Phe Ile Cys Ser Glu Asn Arg
	20	25 30
Tyr	Ser	Leu Gly Leu Glu Pro Asn Pro Ile Arg Glu Pro Val Ala Phe
	35	40 45
Lys	Val	Arg Lys Ala Ile Val Asp Gly Ile Arg Phe Ser Tyr Lys Lys
	50	55 60
Asp	Gly	Ser Val Trp Leu Gln Asn Arg Met Lys Tyr Pro Val Phe Val
	65	70 75 80
Thr	Ser	Gly Tyr Leu Asp Glu Gln Ser Gly Gly Leu Lys Lys Asp Lys
	85	90 95
Val	His	Lys Val Tyr Gly Cys Ala Ser Ile Lys Thr Phe Gly Phe Asn
	100	105 110
Val	Ser	Lys Gln Ile Ile Arg Asp Ala Leu Leu Ser Lys Gln Met Ala
	115	120 125
Thr	Met	Tyr Leu Gln Gly Lys Leu Thr Pro Met Asn Tyr Ile Tyr Glu
	130	135 140
Lys	Lys	Thr Gln Glu Glu Leu Arg Arg Glu Ala Thr Arg Thr Thr Asp
	145	150 155 160

Ser	Leu	Ala	Lys	Tyr	Cys	Cys	Val	Arg	Val	Ser	Phe	Cys	Lys	Gly	Phe
				165					170						175
Gly	Glu	Ala	Tyr	Pro	Glu	Arg	Pro	Ser	Ile	His	Asp	Cys	Pro	Val	Trp
				180				185							190
Ile	Glu	Leu	Lys	Ile	Asn	Ile	Ala	Tyr	Asp	Phe	Met	Asp			
				195			200								205

<210> 114
 <211> 212
 <212> PRT
 <213> Homo sapiens

<400>	114														
Ile	Ala	Tyr	Phe	Glu	Met	Asp	Val	Gln	Val	Gly	Glu	Thr	Phe	Lys	Val
1			5						10						15
Pro	Ser	Ser	Cys	Pro	Ile	Val	Thr	Val	Asp	Gly	Tyr	Val	Asp	Pro	Ser
			20					25							30
Gly	Gly	Asp	Arg	Phe	Cys	Leu	Gly	Gln	Leu	Ser	Asn	Val	His	Arg	Thr
		35				40						45			
Glu	Ala	Ile	Glu	Arg	Ala	Arg	Leu	His	Ile	Gly	Lys	Gly	Val	Gln	Leu
	50				55				60						
Glu	Cys	Lys	Gly	Glu	Gly	Asp	Val	Trp	Val	Arg	Cys	Leu	Ser	Asp	His
	65				70				75						80
Ala	Val	Phe	Val	Gln	Ser	Tyr	Tyr	Leu	Asp	Arg	Glu	Ala	Gly	Arg	Ala
								85		90					95
Pro	Gly	Asp	Ala	Val	His	Lys	Ile	Tyr	Pro	Ser	Ala	Tyr	Ile	Lys	Val
				100				105							110
Phe	Asp	Leu	Arg	Gln	Cys	His	Arg	Gln	Met	Gln	Gln	Gln	Ala	Ala	Thr
		115				120									125
Ala	Gln	Ala	Ala	Ala	Ala	Gln	Ala	Ala	Ala	Val	Ala	Gly	Asn	Ile	
	130					135					140				
Pro	Gly	Pro	Gly	Ser	Val	Gly	Gly	Ile	Ala	Pro	Ala	Ile	Ser	Leu	Ser
	145					150				155					160
Ala	Ala	Ala	Gly	Ile	Gly	Val	Asp	Asp	Leu	Arg	Arg	Leu	Cys	Ile	Leu
			165						170						175
Arg	Met	Ser	Phe	Val	Lys	Gly	Trp	Gly	Pro	Asp	Tyr	Pro	Arg	Gln	Ser
			180					185							190
Ile	Lys	Glu	Thr	Pro	Cys	Trp	Ile	Glu	Ile	His	Leu	His	Arg	Ala	Leu
		195					200								205
Gln	Leu	Leu	Asp												
	210														

<210> 115
 <211> 50
 <212> PRT
 <213> Caenorhabditis elegans

<220>
 <221> VARIANT
 <222> (1)...(50)
 <223> Xaa = Any Amino Acid

<400>	115														
Leu	Cys	Gly	Xaa	Xaa	Leu	Val	Glu	Ala	Leu	Xaa	Xaa	Val	Cys	Gly	Xaa
1			5						10						15
Arg	Gly	Phe	Phe	Tyr	Thr	Pro	Lys	Thr	Arg	Arg	Lys	Arg	Gly	Ile	Val
				20				25							30

Glu Gln Cys Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Gln Leu Glu Xaa Tyr
35 40 45
Cys Asn
50

<210> 116
<211> 39
<212> PRT
<213> *Caenorhabditis elegans*

<400> 116
Leu Cys Gly Arg His Leu Ala Asp Ala Leu Tyr Phe Val Cys Gly Asn
1 5 10 15
Arg Gly Phe Gly Ile Val Glu Glu Cys Cys His Asn Pro Cys Thr Leu
20 25 30
Tyr Gln Leu Glu Asn Tyr Cys
35

<210> 117
<211> 112
<212> PRT
<213> *Caenorhabditis elegans*

<400> 117
Met Asn Ser Val Phe Thr Ile Ile Phe Val Leu Cys Ala Leu Gln Val
1 5 10 15
Ala Ala Ser Phe Arg Gln Ser Phe Gly Pro Ser Met Ser Glu Glu Ser
20 25 30
Ala Ser Met Gln Leu Leu Arg Glu Leu Gln His Asn Met Met Glu Ser
35 40 45
Ala His Arg Pro Met Pro Arg Ala Arg Arg Val Pro Ala Pro Gly Glu
50 55 60
Thr Arg Ala Cys Gly Arg Lys Leu Ile Ser Leu Val Met Ala Val Cys
65 70 75 80
Gly Asp Leu Cys Asn Pro Gln Glu Gly Lys Asp Ile Ala Thr Glu Cys
85 90 95
Cys Gly Asn Gln Cys Ser Asp Asp Tyr Ile Arg Ser Ala Cys Cys Pro
100 105 110

<210> 118
<211> 106
<212> PRT
<213> *Caenorhabditis elegans*

<400> 118
Met Phe Ser Phe Phe Thr Tyr Phe Leu Leu Ser Ala Leu Leu Leu Ser
1 5 10 15
Ala Ser Cys Arg Gln Pro Ser Met Asp Thr Ser Lys Ala Asp Arg Ile
20 25 30
Leu Arg Glu Ile Glu Met Glu Thr Glu Leu Glu Asn Gln Leu Ser Arg
35 40 45
Ala Arg Arg Val Pro Ala Gly Glu Val Arg Ala Cys Gly Arg Arg Leu
50 55 60
Leu Leu Phe Val Trp Ser Thr Cys Gly Glu Pro Cys Thr Pro Gln Glu
65 70 75 80
Asp Met Asp Ile Ala Thr Val Cys Cys Thr Thr Gln Cys Thr Pro Ser

	85	90	95							
Tyr	Ile	Lys	Gln	Ala	Cys	Cys	Pro	Glu	Lys	
	100				105					

<210> 119
<211> 105
<212> PRT
<213> *Caenorhabditis elegans*

<400> 119

Met	Pro	Pro	Ile	Ile	Leu	Val	Phe	Phe	Leu	Val	Leu	Ile	Pro	Ala	Ser
1					5				10					15	
Gln	Gln	Tyr	Pro	Phe	Ser	Leu	Glu	Ser	Leu	Asn	Asp	Gln	Ile	Ile	Asn
					20				25					30	
Glu	Glu	Val	Ile	Glu	Tyr	Met	Leu	Glu	Asn	Ser	Ile	Arg	Ser	Ser	Arg
					35			40				45			
Thr	Arg	Arg	Val	Pro	Asp	Glu	Lys	Lys	Ile	Tyr	Arg	Cys	Gly	Arg	Arg
					50			55			60				
Ile	His	Ser	Tyr	Val	Phe	Ala	Val	Cys	Gly	Lys	Ala	Cys	Glu	Ser	Asn
					65			70			75			80	
Thr	Glu	Val	Asn	Ile	Ala	Ser	Lys	Cys	Cys	Arg	Glu	Glu	Cys	Thr	Asp
					85				90					95	
Asp	Phe	Ile	Arg	Lys	Gln	Cys	Cys	Pro							
					100				105						

<210> 120
<211> 118
<212> PRT
<213> *Caenorhabditis elegans*

<400> 120

Met	Ile	Val	Thr	Leu	Ile	Val	Phe	Leu	Val	Ile	Gly	Leu	Gln	Met	Ala
1					5				10					15	
His	Leu	Ser	Gln	Val	Ser	Gly	Asn	Asn	Glu	Asn	Gly	Phe	Leu	Asn	Pro
					20				25					30	
Phe	Asp	Leu	Ser	Gln	Trp	Ser	Glu	Glu	Ile	Leu	His	Arg	Gln	Tyr	His
					35			40				45			
His	His	His	His	His	His	Gly	Asn	Arg	Ala	Arg	Arg	Arg	Thr	Leu	Glu
					50			55			60				
Thr	Glu	Lys	Ile	Tyr	Arg	Cys	Gly	Arg	Lys	Leu	Tyr	Thr	Asp	Val	Leu
					65			70			75			80	
Ser	Ala	Cys	Asn	Gly	Pro	Cys	Glu	Pro	Gly	Thr	Glu	Gln	Asp	Leu	Ser
					85				90					95	
Lys	Leu	Cys	Cys	Gly	Asn	Gln	Cys	Thr	Phe	Val	Glu	Ile	Arg	Lys	Ala
					100				105					110	
Cys	Cys	Ala	Asp	Lys	Leu										
					115										

<210> 121
<211> 106
<212> PRT
<213> *Caenorhabditis elegans*

<400> 121

Met	Asn	Ala	Ile	Ile	Phe	Cys	Leu	Leu	Phe	Thr	Thr	Val	Thr	Ala	Thr
1					5				10					15	

Tyr Glu Val Phe Gly Lys Gly Ile Glu His Arg Asn Glu His Leu Ile
20 25 30
Ile Asn Gln Leu Asp Ile Ile Pro Val Glu Ser Thr Pro Thr Pro Asn
35 40 45
Arg Ala Ser Arg Val Gln Lys Arg Leu Cys Gly Arg Arg Leu Ile Leu
50 55 60
Phe Met Leu Ala Thr Cys Gly Glu Cys Asp Thr Asp Ser Ser Glu Asp
65 70 75 80
Leu Ser His Ile Cys Cys Ile Lys Gln Cys Asp Val Gln Asp Ile Ile
85 90 95
Arg Val Cys Cys Pro Asn Ser Phe Arg Lys
100 105

<210> 122
<211> 107
<212> PRT
<213> *Caenorhabditis elegans*

<400> 122
Met Lys Leu Ser Val Val Leu Ala Leu Phe Ile Ile Phe Gln Leu Gly
1 5 10 15
Ala Ala Ser Leu Met Arg Asn Trp Met Phe Asp Phe Glu Lys Glu Leu
20 25 30
Glu His Asp Tyr Asp Asp Ser Glu Ile Gly Phe His Asn Ile His Ser
35 40 45
Leu Met Ala Arg Ser Arg Arg Gly Asp Lys Val Lys Ile Cys Gly Thr
50 55 60
Lys Val Leu Lys Met Val Met Val Met Cys Gly Gly Glu Cys Ser Ser
65 70 75 80
Thr Asn Glu Asn Ile Ala Thr Glu Cys Cys Glu Lys Met Cys Thr Met
85 90 95
Glu Asp Ile Thr Thr Lys Cys Cys Pro Ser Arg
100 105

<210> 123
<211> 73
<212> PRT
<213> *Caenorhabditis elegans*

<400> 123
Met Lys Leu Leu His Ile Phe Ile Ile Phe Leu Leu Phe Gln Ser Cys
1 5 10 15
Ser Asn Lys Met Cys Gln Tyr Ser Lys Lys Lys Tyr Lys Ile Cys Gly
20 25 30
Val Arg Ala Leu Lys His Met Lys Val Tyr Cys Thr Arg Gly Met Thr
35 40 45
Arg Asp Tyr Gly Lys Leu Leu Val Thr Cys Cys Ser Lys Gly Cys Asn
50 55 60
Ala Ile Asp Ile Gln Arg Ile Cys Leu
65 70

<210> 124
<211> 109
<212> PRT
<213> *Caenorhabditis elegans*

<400> 124
Met Tyr Trp Phe Arg Gln Val Tyr Arg Pro Ser Phe Phe Phe Gly Phe
1 5 10 15
Leu Ala Ile Leu Leu Leu Ser Ser Pro Thr Pro Ser Asp Ala Ser Ile
20 25 30
Arg Leu Cys Gly Ser Arg Leu Thr Thr Leu Leu Ala Val Cys Arg
35 40 45
Asn Gln Leu Cys Thr Gly Leu Thr Ala Phe Lys Arg Ser Ala Asp Gln
50 55 60
Ser Tyr Ala Pro Thr Thr Arg Asp Leu Phe His Ile His His Gln Gln
65 70 75 80
Lys Arg Gly Gly Ile Ala Thr Glu Cys Cys Glu Lys Arg Cys Ser Phe
85 90 95
Ala Tyr Leu Lys Thr Phe Cys Cys Asn Gln Asp Asp Asn
100 105

<210> 125
<211> 110
<212> PRT
<213> Homo sapiens

<400> 125
Met Ala Leu Trp Met Arg Leu Leu Pro Leu Leu Ala Leu Leu Ala Leu
1 5 10 15
Trp Gly Pro Asp Pro Ala Ala Ala Phe Val Asn Gln His Leu Cys Gly
20 25 30
Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu Arg Gly Phe
35 40 45
Phe Tyr Thr Pro Lys Thr Arg Arg Glu Ala Glu Asp Leu Gln Val Gly
50 55 60
Gln Val Glu Leu Gly Gly Pro Gly Ala Gly Ser Leu Gln Pro Leu
65 70 75 80
Ala Leu Glu Gly Ser Leu Gln Lys Arg Gly Ile Val Glu Gln Cys Cys
85 90 95
Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys Asn
100 105 110

<210> 126
<211> 46
<212> PRT
<213> Caenorhabditis elegans

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 126
Ala Cys Gly Arg Arg Leu Leu Leu Phe Val Trp Ser Thr Cys Gly Glu
1 5 10 15
Pro Cys Thr Xaa Xaa Xaa Gln Glu Asp Met Asp Ile Ala Thr Val Cys
20 25 30
Cys Thr Thr Gln Cys Thr Pro Ser Tyr Ile Lys Gln Ala Cys
35 40 45

<210> 127

<211> 46
<212> PRT
<213> *Caenorhabditis elegans*

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 127
Ala Cys Gly Arg Lys Leu Ile Ser Leu Val Met Ala Val Cys Gly Asp
1 5 10 15
Leu Cys Asn Xaa Xaa Xaa Gln Glu Gly Lys Asp Ile Ala Thr Glu Cys
20 25 30
Cys Gly Asn Gln Cys Ser Asp Asp Tyr Ile Arg Ser Ala Cys
35 40 45

<210> 128
<211> 46
<212> PRT
<213> *Caenorhabditis elegans*

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 128
Arg Cys Gly Arg Arg Ile His Ser Tyr Val Phe Ala Val Cys Gly Lys
1 5 10 15
Ala Cys Glu Xaa Xaa Xaa Ser Thr Glu Val Asn Ile Ala Ser Lys Cys
20 25 30
Cys Arg Glu Glu Cys Thr Asp Asp Phe Ile Arg Lys Gln Cys
35 40 45

<210> 129
<211> 46
<212> PRT
<213> *Caenorhabditis elegans*

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 129
Arg Cys Gly Arg Lys Leu Tyr Thr Asp Val Leu Ser Ala Cys Asn Gly
1 5 10 15
Pro Cys Glu Xaa Xaa Xaa Gly Thr Glu Gln Asp Leu Ser Lys Leu Cys
20 25 30
Cys Gly Asn Gln Cys Thr Phe Asx Glu Ile Arg Lys Ala Cys
35 40 45

<210> 130
<211> 46
<212> PRT

<213> *Caenorhabditis elegans*

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 130

Ile	Cys	Gly	Thr	Lys	Asx	Leu	Lys	Met	Val	Met	Val	Met	Cys	Gly	Gly
1														15	
Glu	Cys	Ser	Xaa	Xaa	Xaa	Ser	Thr	Asn	Glu	Asn	Ile	Ala	Thr	Glu	Cys
			20											30	
Cys	Glu	Lys	Met	Cys	Thr	Met	Glu	Asp	Ile	Thr	Thr	Lys	Cys		
			35											45	

<210> 131

<211> 46

<212> PRT

<213> *Caenorhabditis elegans*

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 131

Leu	Cys	Gly	Arg	Arg	Leu	Ile	Leu	Phe	Met	Leu	Ala	Thr	Cys	Gly	Glu
1														15	
Cys	Asp	Thr	Xaa	Xaa	Xaa	Asp	Ser	Ser	Glu	Asp	Leu	Ser	His	Ile	Cys
			20											30	
Cys	Ile	Lys	Gln	Cys	Asp	Val	Gln	Asp	Ile	Ile	Arg	Val	Cys		
			35											45	

<210> 132

<211> 46

<212> PRT

<213> *Caenorhabditis elegans*

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 132

Leu	Cys	Gly	Ser	His	Leu	Val	Glu	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu
1														15	
Arg	Gly	Phe	Xaa	Xaa	Xaa	Leu	Gln	Lys	Arg	Gly	Ile	Val	Glu	Gln	Cys
			20											30	
Cys	Thr	Ser	Ile	Cys	Ser	Leu	Tyr	Gln	Leu	Glu	Asn	Tyr	Cys		
			35											45	

<210> 133

<211> 46

<212> PRT

<213> Rabbit

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 133
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Glu
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Thr Pro Lys Ser Gly Ile Val Glu Gln Cys
20 25 30
Cys Thr Ser Ile Cys Ser Leu Tyr Gln Leu Glu Asn Tyr Cys
35 40 45

<210> 134
<211> 46
<212> PRT
<213> Xenopus laevis

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 134
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Asp
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Lys Met Lys Arg Gly Ile Val Glu Gln Cys
20 25 30
Cys His Ser Thr Cys Ser Leu Phe Gln Leu Glu Ser Tyr Cys
35 40 45

<210> 135
<211> 46
<212> PRT
<213> Xenopus laevis

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 135
Leu Cys Gly Ser His Leu Val Glu Ala Leu Tyr Leu Val Cys Gly Asp
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Lys Met Lys Arg Gly Ile Val Glu Gln Cys
20 25 30
Cys His Ser Thr Cys Ser Leu Phe Gln Leu Glu Asn Tyr Cys
35 40 45

<210> 136
<211> 46
<212> PRT
<213> Alligator

<220>
<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 136

Leu	Cys	Gly	Ser	His	Leu	Val	Asp	Ala	Leu	Tyr	Leu	Val	Cys	Gly	Glu
1				5					10				15		
Arg	Gly	Phe	Xaa	Xaa	Xaa	Ser	Pro	Lys	Gly	Gly	Ile	Val	Glu	Gln	Cys
		20						25					30		
Cys	His	Asn	Thr	Cys	Ser	Leu	Tyr	Gln	Leu	Glu	Asn	Tyr	Cys		
		35					40					45			

<210> 137

<211> 46

<212> PRT

<213> Elephant fish

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 137

Leu	Cys	Gly	Ser	His	Leu	Val	Asp	Ala	Leu	Tyr	Phe	Val	Cys	Gly	Glu
1				5					10				15		
Arg	Gly	Phe	Xaa	Xaa	Xaa	Pro	Lys	Gln	Ile	Gly	Ile	Val	Glu	Gln	Cys
		20						25					30		
Cys	His	Asn	Thr	Cys	Ser	Leu	Val	Asn	Leu	Glu	Gly	Tyr	Cys		
		35					40					45			

<210> 138

<211> 46

<212> PRT

<213> Bos taurus

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 138

Leu	Cys	Gly	Ala	Glu	Leu	Val	Asp	Ala	Leu	Gln	Phe	Val	Cys	Gly	Asp
1				5					10				15		
Arg	Gly	Phe	Xaa	Xaa	Xaa	Ala	Pro	Gln	Thr	Gly	Ile	Val	Asp	Glu	Cys
		20						25					30		
Cys	Phe	Arg	Ser	Cys	Asp	Leu	Arg	Arg	Leu	Glu	Met	Tyr	Cys		
		35					40					45			

<210> 139

<211> 46

<212> PRT

<213> Canis

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 139
Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln Phe Val Cys Gly Asp
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Ala Pro Gln Thr Gly Ile Val Asp Glu Cys
20 25 30
Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr Cys
35 40 45

<210> 140

<211> 46

<212> PRT

<213> Horse

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 140

Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Arg Arg Ser Arg Gly Ile Val Glu Glu Cys
20 25 30
Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys
35 40 45

<210> 141

<211> 46

<212> PRT

<213> Homo sapiens

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 141

Leu Cys Gly Gly Glu Leu Val Asp Thr Leu Gln Phe Val Cys Gly Asp
1 5 10 15
Arg Gly Phe Xaa Xaa Xaa Arg Arg Ser Arg Gly Ile Val Glu Glu Cys
20 25 30
Cys Phe Arg Ser Cys Asp Leu Ala Leu Leu Glu Thr Tyr Cys
35 40 45

<210> 142

<211> 46

<212> PRT

<213> Amphioxus

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 142

Leu Cys Gly Ser Thr Leu Ala Asp Val Leu Ser Phe Val Cys Gly Asn

1	5	10	15												
Arg	Gly	Tyr	Xaa	Xaa	Xaa	Arg	Arg	Arg	Arg	Gly	Leu	Val	Glu	Glu	Cys
			20				25				30				
Cys	Tyr	Asn	Val	Cys	Asp	Tyr	Ser	Gln	Leu	Glu	Ser	Tyr	Cys		
			35				40				45				

<210> 143

<211> 46

<212> PRT

<213> Locust

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 143

Tyr	Cys	Gly	Glu	Lys	Leu	Ser	Asn	Ala	Leu	Lys	Leu	Val	Cys	Arg	Gly
1				5					10				15		
Asn	Tyr	Asn	Xaa	Xaa	Xaa	Arg	Arg	Thr	Arg	Gly	Val	Phe	Asp	Glu	Cys
			20					25				30			
Cys	Arg	Lys	Ser	Cys	Ser	Ile	Ser	Glu	Leu	Gln	Thr	Tyr	Cys		
			35				40				45				

<210> 144

<211> 46

<212> PRT

<213> Bommo

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 144

Tyr	Cys	Gly	Arg	His	Leu	Ala	Arg	Thr	Leu	Ala	Asp	Leu	Cys	Trp	Glu
1				5					10				15		
Ala	Gly	Val	Xaa	Xaa	Xaa	Arg	Gly	Lys	Arg	Gly	Ile	Val	Asp	Glu	Cys
			20					25				30			
Cys	Leu	Arg	Pro	Cys	Ser	Val	Asp	Val	Leu	Leu	Ser	Tyr	Cys		
			35				40				45				

<210> 145

<211> 46

<212> PRT

<213> Bommo

<220>

<221> VARIANT

<222> (1)...(46)

<223> Xaa = Any Amino Acid

<400> 145

Tyr	Cys	Gly	Arg	His	Leu	Ala	Asp	Thr	Leu	Ala	Asp	Leu	Cys	Phe	Gly
1				5					10				15		
Val	Glu	Lys	Xaa	Xaa	Xaa	Arg	Gly	Lys	Arg	Gly	Val	Val	Asp	Glu	Cys

	20	25	30										
Cys	Phe	Arg	Pro	Cys	Thr	Leu	Asp	Val	Leu	Leu	Ser	Tyr	Cys
	35		40									45	

<210> 146
<211> 46
<212> PRT
<213> Horn worm

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 146
Ile Cys Gly Arg His Leu Ala Arg Thr Leu Ala Asp Leu Cys Pro Asn
1 5 10 15
Val Glu Tyr Xaa Xaa Xaa Gly Lys Arg Ala Gly Val Ala Asp Asp Cys
20 25 30
Cys Asx Asn Ser Cys Thr Met Asp Val Leu Leu Ser Tyr Cys
35 40 45

<210> 147
<211> 46
<212> PRT
<213> Bombyx mori

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 147
Tyr Cys Gly Arg Arg Leu Ala Thr Met Leu Ser Phe Val Cys Asp Asn
1 5 10 15
Gln Tyr Gln Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Ala Glu Glu Cys
20 25 30
Cys Asn Lys Pro Cys Thr Glu Asn Glu Leu Leu Gly Tyr Cys
35 40 45

<210> 148
<211> 46
<212> PRT
<213> Bombyx mori

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 148
Tyr Cys Gly Arg Arg Leu Ala Thr Met Leu Leu Tyr Val Cys Asp Asn
1 5 10 15
Gln Tyr Gln Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Val Glu Glu Cys
20 25 30
Cys Asn Lys Pro Cys Thr Glu Asn Glu Leu Leu Gly Tyr Cys

35

40

45

<210> 149
<211> 46
<212> PRT
<213> Bombyx mori

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 149
Tyr Cys Gly Arg Arg Leu Ala Ile Met Leu Ser Tyr Leu Cys Asp Asn
1 5 10 15
Gln Tyr Leu Xaa Xaa Xaa Gly Lys Arg Gln Gly Ile Ala Glu Glu Cys
20 25 30
Cys Asn Lys Pro Cys Thr Glu Asp Glu Leu Leu Gly Tyr Cys
35 40 45

<210> 150
<211> 46
<212> PRT
<213> Caenorhabditis elegans

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 150
Leu Cys Gly Ser Arg Leu Thr Thr Leu Leu Ala Val Cys Arg Asn
1 5 10 15
Gln Leu Cys Xaa Xaa Xaa Gln Lys Arg Gly Gly Ile Ala Thr Glu Cys
20 25 30
Cys Glu Lys Arg Cys Ser Phe Ala Tyr Leu Lys Thr Phe Cys
35 40 45

<210> 151
<211> 46
<212> PRT
<213> Moi 3

<220>
<221> VARIANT
<222> (1)...(46)
<223> Xaa = Any Amino Acid

<400> 151
Leu Cys Gly Ser Thr Leu Ala Asn Met Val Gln Trp Leu Cys Ser Thr
1 5 10 15
Tyr Thr Thr Xaa Xaa Xaa Glu Ser Arg Pro Ser Ile Val Cys Glu Cys
20 25 30
Cys Phe Asn Gln Cys Thr Val Gln Glu Leu Leu Ala Tyr Cys
35 40 45

<210> 152
 <211> 46
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> (1)...(46)
 <223> Xaa = Any Amino Acid

<400> 152
 Leu Cys Gly Arg Glu Leu Val Arg Ala Gln Ile Ala Ile Cys Gly Met
 1 5 10 15
 Ser Thr Trp Xaa Xaa Xaa Arg Pro Tyr Val Ala Leu Phe Glu Lys Cys
 20 25 30
 Cys Leu Ile Gly Cys Thr Lys Arg Ser Leu Ala Lys Tyr Cys
 35 40 45

<210> 153
 <211> 46
 <212> PRT
 <213> Homo sapiens

<220>
 <221> VARIANT
 <222> (1)...(46)
 <223> Xaa = Any Amino Acid

<400> 153
 Leu Cys Gly His His Phe Val Arg Ala Leu Val Arg Val Cys Gly Gly
 1 5 10 15
 Pro Arg Trp Xaa Xaa Xaa Ala Ala Ala Thr Asn Pro Ala Arg Tyr Cys
 20 25 30
 Cys Leu Ser Gly Cys Thr Gln Gln Asp Leu Leu Thr Leu Cys
 35 40 45

<210> 154
 <211> 541
 <212> PRT
 <213> Caenorhabditis elegans

<400> 154
 Met Ser Met Thr Ser Leu Ser Thr Lys Ser Arg Arg Gln Glu Asp Val
 1 5 10 15
 Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn Trp
 20 25 30
 Arg Pro Arg Tyr Phe Met Ile Phe Asn Asp Gly Ala Leu Leu Gly Phe
 35 40 45
 Arg Ala Lys Pro Lys Glu Gly Gln Pro Phe Pro Glu Pro Leu Asn Asp
 50 55 60
 Phe Met Ile Lys Asp Ala Ala Thr Met Leu Phe Glu Lys Pro Arg Pro
 65 70 75 80
 Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile Glu Arg
 85 90 95
 Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile His Ala
 100 105 110
 Ile Glu Ser Ile Ser Lys Lys Tyr Lys Gly Thr Asn Ala Asn Pro Gln

115	120	125
Glu Glu Leu Met Glu Thr Asn Gln Gln Pro Lys Ile Asp Glu Asp Ser		
130	135	140
Glu Phe Ala Gly Ala Ala His Ala Ile Met Gly Gln Pro Ser Ser Gly		
145	150	155
His Gly Asp Asn Cys Ser Ile Asp Phe Arg Ala Ser Met Ile Ser Ile		160
165	170	175
Ala Asp Thr Ser Glu Ala Ala Lys Arg Asp Lys Ile Thr Met Glu Asp		
180	185	190
Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe Gly Lys Val Ile		
195	200	205
Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala Ile Lys Ile Leu		
210	215	220
Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala His Thr Leu Thr		
225	230	235
Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe Leu Thr Glu Leu		240
245	250	255
Lys Tyr Ser Phe Gln Glu Gln His Tyr Leu Cys Phe Val Met Gln Phe		
260	265	270
Ala Asn Gly Gly Glu Leu Phe Thr His Val Arg Lys Cys Gly Thr Phe		
275	280	285
Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ala Glu Ile Val Leu Ala Leu		
290	295	300
Gly Tyr Leu His Arg Cys Asp Ile Val Tyr Arg Asp Met Lys Leu Glu		
305	310	315
Asn Leu Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala Asp Phe Gly		320
325	330	335
Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser Thr Phe Cys		
340	345	350
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu Asp Asp His Asp Tyr		
355	360	365
Gly Arg Cys Val Asp Trp Trp Gly Val Gly Val Val Met Tyr Glu Met		
370	375	380
Met Cys Gly Arg Leu Pro Phe Tyr Ser Lys Asp His Asn Lys Leu Phe		
385	390	395
Glu Leu Ile Met Ala Gly Asp Leu Arg Phe Pro Ser Lys Leu Ser Gln		400
405	410	415
Glu Ala Arg Thr Leu Leu Thr Gly Leu Leu Val Lys Asp Pro Thr Gln		
420	425	430
Arg Leu Gly Gly Pro Glu Asp Ala Leu Glu Ile Cys Arg Ala Asp		
435	440	445
Phe Phe Arg Thr Val Asp Trp Glu Ala Thr Tyr Arg Lys Glu Ile Glu		
450	455	460
Pro Pro Tyr Lys Pro Asn Val Gln Ser Glu Thr Asp Thr Ser Tyr Phe		
465	470	475
Asp Asn Glu Phe Thr Ser Gln Pro Val Gln Leu Thr Pro Pro Ser Arg		480
485	490	495
Ser Gly Ala Leu Ala Thr Val Asp Glu Gln Glu Glu Met Gln Ser Asn		
500	505	510
Phe Thr Gln Phe Ser Phe His Asn Val Met Gly Ser Ile Asn Arg Ile		
515	520	525
His Glu Ala Ser Glu Asp Asn Glu Asp Tyr Asp Met Gly		
530	535	540

<210> 155

<211> 546

<212> PRT

<213> *Caenorhabditis elegans*

<400> 155

Met Ser Met Thr Ser Leu Ser Thr Lys Ser Arg Arg Gln Glu Asp Val
1 5 10 15
Val Ile Glu Gly Trp Leu His Lys Lys Gly Glu His Ile Arg Asn Trp
20 25 30
Arg Pro Arg Tyr Phe Met Ile Phe Asn Asp Gly Ala Leu Leu Gly Phe
35 40 45
Arg Ala Lys Pro Lys Glu Gly Gln Pro Phe Pro Glu Pro Leu Asn Asp
50 55 60
Phe Met Ile Lys Asp Ala Ala Thr Met Leu Phe Glu Lys Pro Arg Pro
65 70 75 80
Asn Met Phe Met Val Arg Cys Leu Gln Trp Thr Thr Val Ile Glu Arg
85 90 95
Thr Phe Tyr Ala Glu Ser Ala Glu Val Arg Gln Arg Trp Ile His Ala
100 105 110
Ile Glu Ser Ile Ser Lys Lys Tyr Lys Gly Thr Asn Ala Asn Pro Gln
115 120 125
Glu Glu Leu Met Glu Thr Asn Gln Gln Pro Lys Ile Asp Glu Asp Ser
130 135 140
Glu Phe Ala Gly Ala Ala His Ala Ile Met Gly Gln Pro Ser Ser Gly
145 150 155 160
His Gly Asp Asn Cys Ser Ile Asp Phe Arg Ala Ser Met Ile Ser Ile
165 170 175
Ala Asp Thr Ser Glu Ala Ala Lys Arg Asp Lys Ile Thr Met Glu Asp
180 185 190
Phe Asp Phe Leu Lys Val Leu Gly Lys Gly Thr Phe Gly Lys Val Ile
195 200 205
Leu Cys Lys Glu Lys Arg Thr Gln Lys Leu Tyr Ala Ile Lys Ile Leu
210 215 220
Lys Lys Asp Val Ile Ile Ala Arg Glu Glu Val Ala His Thr Leu Thr
225 230 235 240
Glu Asn Arg Val Leu Gln Arg Cys Lys His Pro Phe Leu Thr Glu Leu
245 250 255
Lys Tyr Ser Phe Gln Thr Asn Asp Arg Leu Cys Phe Val Met Glu Phe
260 265 270
Ala Ile Gly Gly Asp Leu Tyr Tyr His Leu Asn Arg Glu Val Gln Met
275 280 285
Asn Lys Glu Gly Phe Ser Glu Pro Arg Ala Arg Phe Tyr Gly Ser Glu
290 295 300
Ile Val Leu Ala Leu Gly Tyr Leu His Ala Asn Ser Ile Val Tyr Arg
305 310 315 320
Asp Leu Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys
325 330 335
Ile Ala Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys
340 345 350
Thr Ser Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu
355 360 365
Asp Asp His Asp Tyr Gly Arg Cys Val Asp Trp Trp Gly Val Gly Val
370 375 380
Val Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Ser Lys Asp
385 390 395 400
His Asn Lys Leu Phe Glu Leu Ile Met Ala Gly Asp Leu Arg Phe Pro
405 410 415
Ser Lys Leu Ser Gln Glu Ala Arg Thr Leu Leu Thr Gly Leu Leu Val
420 425 430
Lys Asp Pro Thr Gln Arg Leu Gly Gly Pro Glu Asp Ala Leu Glu
435 440 445
Ile Cys Arg Ala Asp Phe Phe Arg Thr Val Asp Trp Glu Ala Thr Tyr
450 455 460

Arg	Lys	Glu	Ile	Glu	Pro	Pro	Tyr	Lys	Pro	Asn	Val	Gln	Ser	Glu	Thr
465				470				475						480	
Asp	Thr	Ser	Tyr	Phe	Asp	Asn	Glu	Phe	Thr	Ser	Gln	Pro	Val	Gln	Leu
			485					490						495	
Thr	Pro	Pro	Ser	Arg	Ser	Gly	Ala	Leu	Ala	Thr	Val	Asp	Glu	Gln	Glu
			500					505					510		
Glu	Met	Gln	Ser	Asn	Phe	Thr	Gln	Phe	Ser	Phe	His	Asn	Val	Met	Gly
		515					520					525			
Ser	Ile	Asn	Arg	Ile	His	Glu	Ala	Ser	Glu	Asp	Asn	Glu	Asp	Tyr	Asp
	530					535					540				
Met	Gly														
	545														

<210> 156

<211> 483

<212> PRT

<213> *Caenorhabditis elegans*

<400> 156

Met	Ser	Thr	Glu	Asn	Ala	His	Leu	Gln	Lys	Glu	Asp	Ile	Val	Ile	Glu
1					5				10					15	
Ser	Trp	Leu	His	Lys	Lys	Gly	Glu	His	Ile	Arg	Asn	Trp	Arg	Pro	Arg
					20				25					30	
Tyr	Phe	Ile	Leu	Phe	Arg	Asp	Gly	Thr	Leu	Leu	Gly	Phe	Arg	Ser	Lys
					35				40					45	
Pro	Lys	Glu	Asp	Gln	Pro	Leu	Pro	Glu	Pro	Leu	Asn	Asn	Phe	Met	Ile
					50				55					60	
Arg	Asp	Ala	Ala	Thr	Val	Cys	Leu	Asp	Lys	Pro	Arg	Pro	Asn	Met	Phe
					65				70					80	
Ile	Val	Arg	Cys	Leu	Gln	Trp	Thr	Thr	Val	Ile	Glu	Arg	Thr	Phe	Tyr
					85				90					95	
Ala	Asp	Ser	Ala	Asp	Phe	Arg	Gln	Met	Trp	Ile	Glu	Ala	Ile	Gln	Ala
					100				105					110	
Val	Ser	Ser	His	Asn	Arg	Leu	Lys	Glu	Asn	Ala	Gly	Asn	Thr	Ser	Met
					115				120					125	
Gln	Glu	Glu	Asp	Thr	Asn	Gly	Asn	Pro	Ser	Gly	Glu	Ser	Asp	Val	Asn
					130				135					140	
Met	Asp	Ala	Thr	Ser	Thr	Arg	Ser	Asp	Asn	Asp	Phe	Glu	Ser	Thr	Val
					145				150					160	
Met	Asn	Ile	Asp	Glu	Pro	Glu	Glu	Val	Pro	Arg	Lys	Asn	Thr	Val	Thr
					165				170					175	
Met	Asp	Asp	Phe	Asp	Phe	Leu	Lys	Val	Leu	Gly	Gln	Gly	Thr	Phe	Gly
					180				185					190	
Lys	Val	Ile	Leu	Cys	Arg	Glu	Lys	Ser	Ser	Asp	Lys	Leu	Tyr	Ala	Ile
					195				200					205	
Lys	Ile	Ile	Arg	Lys	Glu	Met	Val	Val	Asp	Arg	Ser	Glu	Val	Ala	His
					210				215					220	
Thr	Leu	Thr	Glu	Asn	Arg	Val	Leu	Tyr	Ala	Cys	Val	His	Pro	Phe	Leu
					225				230					240	
Thr	Leu	Leu	Lys	Tyr	Ser	Phe	Gln	Ala	Gln	Tyr	His	Ile	Cys	Phe	Val
					245				250					255	
Met	Glu	Phe	Ala	Asn	Gly	Gly	Glu	Leu	Phe	Thr	His	Leu	Gln	Arg	Cys
					260				265					270	
Lys	Thr	Phe	Ser	Glu	Ala	Arg	Thr	Arg	Phe	Tyr	Gly	Ser	Glu	Ile	Ile
					275				280					285	
Leu	Ala	Leu	Gly	Tyr	Leu	His	His	Arg	Asn	Ile	Val	Tyr	Arg	Asp	Met
					290				295					300	
Lys	Leu	Glu	Asn	Leu	Leu	Leu	Asp	Arg	Asp	Gly	His	Ile	Lys	Ile	Thr

305	310	315	320
Asp Phe Gly Leu Cys Lys Glu Glu Ile Lys Tyr Gly Asp Lys Thr Ser			
325	330	335	
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Ile Glu Asp			
340	345	350	
Ile Asp Tyr Asp Arg Ser Val Asp Trp Trp Gly Val Gly Val Val Met			
355	360	365	
Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Ser Ala Lys Glu Asn Gly			
370	375	380	
Lys Leu Phe Glu Leu Ile Thr Thr Cys Asp Leu Lys Phe Pro Asn Arg			
385	390	395	400
Leu Ser Pro Glu Ala Val Thr Leu Leu Ser Gly Leu Leu Glu Arg Val			
405	410	415	
Pro Ala Lys Arg Leu Gly Ala Gly Pro Asp Asp Ala Arg Glu Val Ser			
420	425	430	
Arg Ala Glu Phe Phe Lys Asp Val Asp Trp Glu Ala Thr Leu Arg Lys			
435	440	445	
Glu Val Glu Pro Pro Phe Lys Pro Asn Val Met Ser Glu Thr Asp Thr			
450	455	460	
Ser Phe Phe Asp Arg Val Arg Tyr Val Ser Ile Leu Leu Lys Val Ser			
465	470	475	480
Glu Ala Ile			

<210> 157
 <211> 480
 <212> PRT
 <213> Homo sapiens

<400> 157			
Met Ser Asp Val Ala Ile Val Lys Glu Gly Trp Leu His Lys Arg Gly			
1	5	10	15
Glu Tyr Ile Lys Thr Trp Arg Pro Arg Tyr Phe Leu Leu Lys Asn Asp			
20	25	30	
Gly Thr Phe Ile Gly Tyr Lys Glu Arg Pro Gln Val Asp Val Gln Arg			
35	40	45	
Glu Ala Pro Leu Asn Asn Phe Ser Val Ala Gln Cys Gln Leu Met Lys			
50	55	60	
Thr Glu Arg Pro Arg Pro Asn Thr Phe Ile Ile Arg Cys Leu Gln Trp			
65	70	75	80
Thr Thr Val Ile Glu Arg Thr Phe His Val Glu Thr Pro Glu Glu Arg			
85	90	95	
Glu Glu Trp Thr Thr Ala Ile Gln Thr Val Ala Asp Gly Leu Lys Lys			
100	105	110	
Gln Glu Glu Glu Glu Met Asp Phe Arg Ser Gly Ser Pro Ser Asp Asn			
115	120	125	
Ser Gly Ala Glu Glu Met Glu Val Ser Leu Ala Lys Pro Lys His Arg			
130	135	140	
Val Thr Met Asn Glu Phe Glu Tyr Leu Lys Leu Leu Gly Lys Gly Thr			
145	150	155	160
Phe Gly Lys Val Ile Leu Val Lys Glu Lys Ala Thr Gly Arg Tyr Tyr			
165	170	175	
Ala Met Lys Ile Leu Lys Lys Glu Val Ile Val Ala Lys Asp Glu Val			
180	185	190	
Ala His Thr Leu Thr Glu Asn Arg Val Leu Gln Asn Ser Arg His Pro			
195	200	205	
Phe Leu Thr Ala Leu Lys Tyr Ser Phe Gln Thr His Asp Arg Leu Cys			
210	215	220	

Phe Val Met Glu Tyr Ala Asn Glu Gly Leu Phe Phe His Leu Ser
 225 230 235 240
 Arg Glu Arg Val Phe Ser Glu Asp Phe Ala Phe Arg Tyr Glu Ala Glu
 245 250 255
 Ile Val Ser Ala Leu Asp Tyr Leu His Ser Glu Lys Asn Val Tyr
 260 265 270
 Arg Asp Leu Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Glu His Ile
 275 280 285
 Lys Ile Thr Asp Phe Glu Leu Cys Lys Glu Glu Ile Lys Asp Glu Ala
 290 295 300
 Thr Met Lys Thr Phe Cys Glu Tyr Pro Glu Tyr Leu Ala Pro Glu Val
 305 310 315 320
 Leu Glu Asp Asn Asp Tyr Glu Arg Ala Val Asp Trp Trp Glu Leu Glu
 325 330 335
 Val Val Met Tyr Glu Met Cys Glu Arg Leu Pro Phe Tyr Asn Glu
 340 345 350
 Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile Arg Phe
 355 360 365
 Pro Arg Thr Leu Glu Pro Glu Ala Lys Ser Leu Leu Ser Glu Leu
 370 375 380
 Lys Lys Asp Pro Lys Glu Arg Leu Glu Glu Ser Glu Asp Ala Lys
 385 390 395
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 400 405 410 415
 Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Glu Val Thr Ser Glu
 420 425 430 435
 Thr Asp Thr Arg Tyr Phe Asp Glu Phe Thr Ala Glu Met Ile Thr
 440 445 450 455
 Ile Thr Pro Pro Asp Glu Asp Ser Met Glu Cys Val Asp Ser Glu
 460 465 470 475
 Arg Arg Pro His Phe Pro Glu Phe Ser Tyr Ser Ala Ser Ser Thr Ala
 480

<400> 158
 cataaatc cagtaatgg taataatttc aatttcagat catttcgat gggaggatc
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 120
 gaaaggccgc caacgggttag gaaactatgtt ctagacgaaatc atcggtatcg
 180
 atacccatca actacccatca actacccatca atcggtatcg
 240
 tgggtgcac ttccaaatcg tttccatccatc tccatccatc
 300
 agcccaat tttccatca actacccatca actacccatca atcggtatcg
 360
 cttttttttt tttttttttt tttttttttt tttttttttt
 420
 aatgtatata tttttttttt tttttttttt tttttttttt
 480
 atggcttcctg ttccaaatcg tttccatccatc tccatccatc
 540
 cttttttttt tttttttttt tttttttttt tttttttttt
 600
 tttaggcctt ttttttatata aatgttttgc agttttttgtt tttttttttt
 660
 acgtttttttt tttttttttt tttttttttt tttttttttt
 720
 ttccacccatc cttttttttt tttttttttt tttttttttt
 780
 atatattatc aatattttttt tttttttttt tttttttttt
 840
 ttccacatc tttttttttt tttttttttt tttttttttt
 900
 caaatattttt tttttttttt tttttttttt tttttttttt
 960
 tttcccccac ttccatccatc tttttttttt tttttttttt
 1020
 agacatccatc acgacttcata gttttttttt tttttttttt
 1080
 gttttttttt gttttttttt gttttttttt gttttttttt
 1140

<210> 158
 <211> 6250
 <212> DNA
 <213> Caenorhabditis elegans

Arg Arg Pro His Phe Pro Glu Phe Ser Tyr Ser Ala Ser Ser Thr Ala
 465 470 475 480
 Ile Thr Pro Pro Asp Glu Asp Ser Met Glu Cys Val Asp Ser Glu
 490 495 500 505
 Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Glu Val Thr Ser Glu
 510 515 520 525
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 530 535 540 545
 Lys Lys Asp Pro Lys Glu Arg Leu Glu Glu Ser Glu Asp Ala Lys
 550 555 560 565
 Pro Arg Thr Leu Glu Pro Glu Ala Lys Ser Leu Leu Ser Glu Leu
 570 575 580 585
 Lys Lys Asp Pro Lys Glu Arg Leu Glu Glu Ser Glu Asp Ala Lys
 590 595 600 605
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 610 615 620 625
 Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Glu Val Thr Ser Glu
 630 635 640 645
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 650 655 660 665
 Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Glu Val Thr Ser Glu
 670 675 680 685
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 690 695 700 705
 Tyr Glu Lys Lys Leu Ser Pro Pro Phe Lys Pro Glu Val Thr Ser Glu
 710 715 720 725
 Glu Ile Met Glu His Arg Phe Phe Ala Glu Ile Val Trp Glu His Val
 730 735 740 745
 Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile Arg Phe
 750 755 760 765
 Pro Arg Thr Leu Glu Pro Glu Ala Lys Ser Leu Leu Ser Glu Leu
 770 775 780 785
 Lys Ile Thr Asp Phe Glu Leu Cys Lys Glu Glu Ile Lys Asp Glu Ala
 790 795 800 805
 Thr Met Lys Thr Phe Cys Glu Tyr Pro Glu Tyr Leu Ala Pro Glu Val
 810 815 820 825
 Leu Glu Asp Asn Asp Tyr Glu Arg Ala Val Asp Trp Trp Glu Leu Glu
 830 835 840 845
 Val Val Met Tyr Glu Met Cys Glu Arg Leu Pro Phe Tyr Asn Glu
 850 855 860 865
 Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile Arg Phe
 870 875 880 885
 Pro Arg Thr Leu Glu Pro Glu Ala Lys Ser Leu Leu Ser Glu Leu
 890 895 900 905
 Lys Ile Thr Asp Phe Glu Leu Cys Lys Glu Glu Ile Lys Asp Glu Ala
 910 915 920 925
 Thr Met Lys Thr Phe Cys Glu Tyr Pro Glu Tyr Leu Ala Pro Glu Val
 930 935 940 945
 Leu Glu Asp Asn Asp Tyr Glu Arg Ala Val Asp Trp Trp Glu Leu Glu
 950 955 960 965
 Val Val Met Tyr Glu Met Cys Glu Arg Leu Pro Phe Tyr Asn Glu
 970 975 980 985
 Asp His Glu Lys Leu Phe Glu Leu Ile Leu Met Glu Glu Ile Arg Phe
 990 995 1000 1005
 Pro Arg Thr Leu Glu Pro Glu Ala Lys Ser Leu Leu Ser Glu Leu

Phe Ala Ser Glu Ile Leu Thr Glu Leu Glu Phe Leu His Asp Asn Lys
 165 170 175
 Glu Ser Leu Cys His Phe Glu Ser Phe Asp Met Leu Thr Ser Lys Phe
 145 150 155 160
 Ala Arg Ile Tyr Phe Val Ile Glu Leu Val Glu Asn Glu Asp Leu Glu
 130 135 140
 Glu Gly His Pro Phe Val Thr Glu Leu Tyr Thr His Phe His Asp Glu
 115 120 125
 Ala Ile Ile Arg Glu Lys Asn Ile Leu Thr Tyr Leu Ser Glu Glu Cys
 100 105 110
 Val Lys Val Leu Glu Lys Ser Tyr Leu Asn Arg His Glu Lys Met Asp
 85 90 95
 Ser Glu Val Phe Arg Cys Arg Glu Val Ala Thr Asp Ala Met Phe Ala
 65 70 75
 Thr Ser Asn Asp Phe Met Phe Leu Glu Ser Met Glu Glu Gly Ala Tyr
 50 55 60
 Ala Glu Asp Leu Ile Ala Lys Ser Ile Lys Glu Gly Cys Pro Lys Arg
 35 40 45
 Leu Thr Pro Thr Ala Ser Glu Asn Ser Leu Ser Pro Val Thr
 20 25 30
 Asn Asn Asp Thr Thr Ser Asp Arg Glu Ala Pro Thr Thr Leu Asn
 1 5 10 15
 Met Glu Asp Leu Thr Pro Thr Asn Thr Ser Leu Asp Thr Thr Thr
 <400> 159

<213> Caenorhabditis elegans

<212> PRT
 <211> 632
 <210> 159

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 6180
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 6120
 tcacataccat cttttcttgtt caaaaatgttgc tcaacttgttgc
 6060
 aaggcatttttt gggccacccac gcaatcttttta ttatatttgttgc
 6000
 attttggggatc accctggggatc accatgtttaca ttatattgttgc
 5940
 ttatggatgtt tagacgggttt aatatttttg atgatttttaa ttatatttgttgc
 5880
 taatagggtttt ttatttttccaa tggccatcttca aatattttttt
 5820
 agatttctggaa aatattttccaa tcttcataaca aacatataattt
 5760
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 5700
 aatattttttttt tttttttttt tttttttttt tttttttttt
 5640
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 5580
 caaaaaaaaatg tttttttttt tttttttttt tttttttttt
 5520
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 5460
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 5400
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 5340
 aggccatccatc tttttttttt tttttttttt tttttttttt
 5280
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 5220
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 5160
 acggccgttca tttttttttt tttttttttt tttttttttt
 5100
 gggatatttgg tttttttttt tttttttttt tttttttttt
 5040
 gggatatttgg tttttttttt tttttttttt tttttttttt
 4980
 gggatatttgg tttttttttt tttttttttt tttttttttt
 4920
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 4860
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 4800
 cttttttttttt tttttttttt tttttttttt tttttttttt
 4740

<210> 163
<211> 54

<400> 162
Ser Pro Phe Lys Trp Ser Pro Ser Asp Trp Thr Phe Arg Pro Arg Ser
1 5 10 15
Ser Asn Ala Ser Ser Arg Leu Ser Pro Glu Leu Glu
20 25

<210> 162
<211> 28
<212> PRT
<213> Homo sapiens

<400> 161
Ser Pro Val Gly His Phe Ala Lys Trp Ser Gly Ser Pro Cys Ser Arg
1 5 10 15
Asn Arg Glu Ala Asp Met Trp Thr Thr Phe Arg Pro Arg Ser Ser
20 25 30
Ser Asn Ala Ser Ser Val Ser Thr Arg Leu Ser Pro Leu Arg Pro Glu
35 40
Ser Glu Val Leu Ala Glu
50

<210> 161
<211> 54
<212> PRT
<213> Homo sapiens

<400> 161
Lys Lys Leu Ser Met Glu Met Asp Lys Ser Pro
620 625
Lys Glu Lys Glu Lys Lys Ala Leu Lys Ala Glu Glu Val Ser
640 645
Lys Lys Ser Arg Lys Glu Met Met Arg Glu Glu Lys Ala Leu Arg Arg
595 600
Phe Asn Ser Ala Met Arg Asp Gly Thr Phe Gly Ser Ile Tyr Gly Lys
580 585 590
Ala Ile Asn Asp Val Arg Lys Arg Tyr Ser Val Thr Ile Glu Lys Thr
560 565
Val Tyr Leu Phe Asp Leu Glu Lys Ala Asp Glu Trp Cys Lys
545 550 555
Val Glu Leu Lys Asn Ser Gly Thr Phe Ile His Thr Pro Asn Arg
530 535
Pro Asn Leu Val Leu Lys Glu Val Pro Trp Thr Pro Cys Met Glu
515 520 525
Arg Met Phe Leu Leu Thr Glu Pro His Leu Tyr Ile Asp Val
495 500
Leu Lys Glu Gly Tyr Leu Glu Lys Lys Arg Gly Leu Phe Ala Arg Arg
485 490 495
Glu Arg Val Lys Asn Pro Phe His Ile Phe Thr Asn Asn Ser Leu Ile
465 470 475
Ser Glu Ala Glu Lys Asn Arg Ala Arg Ala Glu Lys Leu Glu
450 455
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
420 425 430 435
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
440 445
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
460 465
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
475 480
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
490 495
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
500 505
Glu His Arg Gly Asp Pro Phe Val Ser Glu Ile Ala Pro Arg Ala Asn
510 515
Pro Asn Leu Val Leu Lys Glu Val Pro Trp Thr Pro Cys Met Glu
525 530
Val Glu Leu Lys Asn Ser Gly Thr Phe Ile His Thr Pro Asn Arg
540 545
Ala Ile Asn Asp Val Arg Lys Arg Tyr Ser Val Thr Ile Glu Lys Thr
560 565
Val Tyr Leu Phe Asp Leu Glu Lys Ala Asp Glu Trp Cys Lys
580 585 590
Phe Asn Ser Ala Met Arg Asp Gly Thr Phe Gly Ser Ile Tyr Gly Lys
595 600
Lys Glu Lys Glu Lys Lys Ala Leu Lys Ala Glu Glu Val Ser
610 615
Ser Asn Ala Ser Ser Val Ser Thr Arg Leu Ser Pro Leu Arg Pro Glu
620 625
Ser Glu Val Leu Ala Glu
630 635

Glù Ala Pro Glù Val Val Glù Ile Asp Pro Asp Phe Glù Pro Leu Pro
<400> 167

<213> Homo sapiens

<212> PRT

<211> 42

<210> 167

Pro Arg Pro Glù Ser Pro
Ala Asp Pro Asp Phe Glù Pro Arg Ser Cys Thr Trp Pro Leu
1 5 10 15
<400> 166

<213> Homo sapiens

<212> PRT

<211> 22

<210> 166

Glù Pro Ser Glù Pro Pro Glù Val Glù Pro
Arg Pro Arg Ser Cys Thr Trp Pro Leu Pro Arg Pro Glù Ile Ala Asn
1 5 10 15 20 25 30
Lys Ala Ala Ala Ile Ile Asp Leu Asp Pro Asp Phe Glù Pro Glù Ser
<400> 165

<213> Homo sapiens

<212> PRT

<211> 42

<210> 165

Ser Phe Arg Pro Arg Thr Glù Ser Asn Leu Ser Ile Pro Gly Ser Ser
1 5 10 15
<400> 164

<213> Caenorhabditis elegans

<212> PRT

<211> 17

<210> 164

Glù Asp Asp Leu Gly Glù
Ser Asn Ala Ser Thr Ile Ser Gly Arg Leu Ser Pro Ile Met Thr Glù
1 5 10 15 20 25 30 35 40 45
Ser Asn Asp Asp Phe Asp Asn Trp Ser Thr Phe Arg Pro Arg Thr Ser
I 5 10 15
Ser Pro Gly Ser Glù Phe Ser Lys Trp Pro Ala Ser Pro Gly Ser His
<400> 163

<213> Homo sapiens

<212> PRT

Asn Tyr Asp His Leu Lys Glu Pro Arg Arg Arg Leu Leu Thr Glu Pro His Leu Tyr Asp Asn
15 10 5 20 25 30
<400> 171

<213> Mus musculus or Homo sapiens or C elegans
<212> PRT
<211> 47
<210> 171

Asn Lys Glu Ala Glu Asn Pro Trp His Glu Phe Val Glu Asn
15 10 5 20 25 30
<400> 170

Asn Leu Ile Leu Lys Met Glu Pro Val Asp Lys Arg Lys Glu Leu Phe
15 10 5 20 25 30
Ala Arg Arg Glu Leu Leu Leu Thr Glu Glu Pro His Leu Tyr Tyr
35 40 45 50
Val Asp Pro Val Asn Lys Val Leu Lys Glu Ile Pro Trp Ser Glu
55
Glu Leu Arg Pro Glu Ala Lys Asn Phe Lys Thr Phe Val His Thr
60 65 70 75 80
<213> Mus musculus or Homo sapiens

<212> PRT
<211> 80
<210> 170

Pro Arg Thr Trp Pro Arg Thr Trp Pro Arg Pro
10 5 1
<400> 169

<213> Caenorhabditis elegans or Homo sapiens
<212> PRT
<211> 14
<210> 169

Thr Phe Met Asn Thr Pro Asp Asp Val Met Met Asn Asp Asp Met Glu
15 10 5 20 25 30
Pro Ile Pro Arg Asp Arg Cys Asn Thr Trp Pro Met Arg Arg Pro Glu
35
Leu Glu Pro Pro Leu Asn Ser Ser Pro
<400> 168

<213> Caenorhabditis elegans
<212> PRT
<211> 41
<210> 168

Arg Pro Arg Ser Ser Cys Thr Trp Pro Leu Pro Arg Pro Glu Phe Ser Glu
15 10 5 20 25 30
Ser Asn Ser Ala Thr Ser Ser Pro Ala Pro
40 35
<400> 167

Val Leu Lys Gly Glu Pro Trp Glu Lys Asn Thr Phe Phe His Thr
35 40 45

<210> 172

<211> 80

<212> PRT

<213> *Caenorhabditis elegans*

<400> 172

Leu Glu Glu Gln Arg Val Lys Asn Pro Phe His Ile Phe Thr Asn Asn
1 5 10 15
Ser Leu Ile Leu Lys Gln Gly Tyr Leu Glu Lys Lys Arg Gly Leu Phe
20 25 30
Ala Arg Arg Arg Met Phe Leu Leu Thr Glu Gly Pro His Leu Leu Tyr
35 40 45
Ile Asp Val Pro Asn Leu Val Leu Lys Gly Glu Val Pro Trp Thr Pro
50 55 60
Cys Met Gln Val Glu Leu Lys Asn Ser Gly Thr Phe Phe Ile His Thr
65 70 75 80

<210> 173

<211> 113

<212> PRT

<213> *Mus musculus* or *Homo sapiens*

<400> 173

Ser Asp Leu Trp Ala Leu Gly Cys Ile Ile Tyr Gln Leu Val Ala Gly
1 5 10 15
Leu Pro Pro Phe Arg Ala Gly Asn Glu Tyr Leu Ile Phe Gln Lys Ile
20 25 30
Ile Lys Leu Glu Tyr Asp Phe Pro Glu Lys Phe Phe Pro Lys Ala Arg
35 40 45
Asp Leu Val Glu Lys Leu Leu Val Leu Asp Ala Thr Lys Arg Leu Gly
50 55 60
Cys Glu Glu Met Glu Gly Tyr Gly Pro Leu Lys Ala His Pro Phe Phe
65 70 75 80
Glu Ser Val Thr Trp Glu Asn Leu His Gln Gln Thr Pro Pro Lys Leu
85 90 95
Thr Ala Tyr Leu Pro Ala Met Ser Glu Asp Asp Glu Asp Cys Tyr Gly
100 105 110
Asn

<210> 174

<211> 48

<212> PRT

<213> *Mus musculus* or *Homo sapiens* or *C elegans*

<400> 174

Asp Trp Leu Gly Cys Ile Gln Ala Gly Pro Pro Phe Arg Ala Asn Tyr
1 5 10 15
Ile Leu Phe Pro Glu Phe Ala Lys Leu Val Leu Glu Pro Leu Ala His
20 25 30
Phe Phe Glu Val Trp Asn Pro Pro Leu Ala Tyr Pro Ala Glu Tyr Asn
35 40 45

<210> 175
<211> 122
<212> PRT
<213> *Caenorhabditis elegans*

<400> 175

Thr	Asp	Ile	Trp	Gly	Leu	Gly	Cys	Ile	Leu	Phe	Gln	Cys	Leu	Ala	Gly
1				5				10					15		
Gln	Pro	Pro	Phe	Arg	Ala	Val	Asn	Gln	Tyr	His	Leu	Leu	Lys	Arg	Ile
				20				25					30		
Gln	Glu	Leu	Asp	Phe	Ser	Phe	Pro	Glu	Gly	Phe	Pro	Glu	Glu	Ala	Ser
	35					40						45			
Glu	Ile	Ile	Ala	Lys	Ile	Leu	Val	Gly	His	Glu	Thr	Leu	Lys	Thr	Glu
	50					55				60					
Tyr	Val	Ile	Phe	Asn	Leu	Gln	Val	Arg	Asp	Pro	Ser	Thr	Arg	Ile	Thr
65					70				75				80		
Ser	Gln	Glu	Leu	Met	Ala	His	Lys	Phe	Phe	Glu	Asn	Val	Asp	Trp	Val
				85					90				95		
Asn	Ile	Ala	Asn	Ile	Lys	Pro	Pro	Val	Leu	His	Ala	Tyr	Ile	Pro	Ala
			100				105						110		
Thr	Phe	Gly	Glu	Pro	Glu	Tyr	Tyr	Ser	Asn						
	115					120									

<210> 176
<211> 72
<212> PRT
<213> *Mus musculus* or *Homo sapiens*

<400> 176

Phe	Gly	Leu	Ser	Tyr	Ala	Lys	Asn	Gly	Glu	Leu	Leu	Lys	Tyr	Ile	Arg
1					5				10				15		
Lys	Ile	Gly	Ser	Phe	Asp	Glu	Thr	Cys	Thr	Arg	Phe	Tyr	Thr	Ala	Glu
					20			25				30			
Ile	Val	Ser	Ala	Leu	Glu	Tyr	Leu	His	Gly	Lys	Gly	Ile	Ile	His	Arg
				35			40				45				
Asp	Leu	Lys	Pro	Glu	Asn	Ile	Leu	Leu	Asn	Glu	Asp	Met	His	Ile	Gln
			50				55				60				
Ile	Thr	Asp	Phe	Gly	Thr	Ala	Lys								
65				70											

<210> 177
<211> 31
<212> PRT
<213> *Mus musculus* or *Homo sapiens* or *C elegans*

<400> 177

Phe	Asn	Gly	Leu	Gly	Ser	Phe	Asp	Phe	Glu	Ile	Leu	Leu	His	Ile	His
1					5				10				15		
Arg	Asp	Lys	Pro	Asn	Leu	Asp	His	Ile	Ile	Thr	Asp	Phe	Gly	Ala	
					20			25				30			

<210> 178
<211> 72
<212> PRT
<213> *Caenorhabditis elegans*

<400> 178
Phe Val Ile Gly Leu Val Glu Asn Gly Asp Leu Gly Glu Ser Leu Cys
1 5 10 15
His Phe Gly Ser Phe Asp Met Leu Thr Ser Lys Phe Phe Ala Ser Glu
20 25 30
Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys Ile Val His Arg
35 40 45
Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp Gly His Ile Leu
50 55 60
Ile Thr Asp Phe Gly Ser Ala Gln
65 70

<210> 179

<211> 48

<212> PRT

<213> Mus musculus or Homo sapiens

<400> 179

Tyr Ala Ile Lys Ile Leu Glu Lys Arg His Ile Ile Lys Glu Asn Lys
1 5 10 15
Val Pro Tyr Val Thr Arg Glu Arg Asp Val Met Ser Arg Leu Asp His
20 25 30
Pro Phe Phe Val Lys Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys Leu
35 40 45

<210> 180

<211> 15

<212> PRT

<213> Mus musculus or Homo sapiens or C elegans

<400> 180

Ala Lys Leu Lys Lys Arg Glu Leu His Pro Phe Leu Tyr Phe Asp
1 5 10 15

<210> 181

<211> 53

<212> PRT

<213> Caenorhabditis elegans

<400> 181

Phe Ala Val Lys Val Leu Gln Lys Ser Tyr Leu Asn Arg His Gln Lys
1 5 10 15
Met Asp Ala Ile Ile Arg Glu Lys Asn Ile Leu Thr Tyr Leu Ser Gln
20 25 30
Glu Cys Gly Gly His Pro Phe Val Thr Gln Leu Tyr Thr His Phe His
35 40 45
Asp Gln Ala Arg Ile
50

<210> 182

<211> 29

<212> PRT

<213> Mus musculus or Homo sapiens

<400> 182

Pro Asn Arg Thr Tyr Tyr Leu Met Asp Pro Ser Gly Asn Ala His Lys
1 5 10 15
Trp Cys Arg Lys Ile Gln Glu Val Trp Arg Gln Arg Tyr
20 25

<210> 183
<211> 15
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans

<400> 183
Pro Asn Arg Tyr Tyr Leu Asp Ala Trp Cys Ile Val Arg Arg Tyr
1 5 10 15

<210> 184
<211> 28
<212> PRT
<213> Caenorhabditis elegans

<400> 184
Pro Asn Arg Val Tyr Tyr Leu Phe Asp Leu Glu Lys Lys Ala Asp Glu
1 5 10 15
Trp Cys Lys Ala Ile Asn Asp Val Arg Lys Arg Tyr
20 25

<210> 185
<211> 25
<212> PRT
<213> Mus musculus or Homo sapiens

<400> 185
Pro Glu Ser Lys Gln Ala Arg Ala Asn Ser Phe Val Gly Thr Ala Gln
1 5 10 15
Tyr Val Ser Pro Glu Leu Leu Thr Glu
20 25

<210> 186
<211> 15
<212> PRT
<213> Mus musculus or Homo sapiens or C elegans

<400> 186
Pro Glu Ala Arg Phe Val Gly Thr Ala Tyr Val Ser Pro Glu Leu
1 5 10 15

<210> 187
<211> 25
<212> PRT
<213> Caenorhabditis elegans

<400> 187
Pro Glu Glu Asn Thr Ala Arg Arg Thr Thr Phe Val Gly Thr Ala Leu
1 5 10 15
Tyr Val Ser Pro Glu Met Leu Ala Asp

<210> 188

<211> 62

<212> PRT

<213> *Caenorhabditis elegans*

<400> 188

Lys	Arg	Thr	Ser	Asn	Asp	Phe	Met	Phe	Leu	Gln	Ser	Met	Gly	Glu	Gly
1															
														15	
Ala	Tyr	Ser	Gln	Val	Phe	Arg	Cys	Arg	Glu	Val	Ala	Thr	Asp	Ala	Met
														30	
Phe	Ala	Val	Lys	Val	Leu	Gln	Lys	Ser	Tyr	Leu	Asn	Arg	His	Gln	Lys
														45	
Met	Asp	Ala	Ile	Ile	Arg	Glu	Lys	Asn	Ile	Leu	Thr	Tyr	Tyr	Leu	
														60	
50															

<210> 189

<211> 21

<212> PRT

<213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 189

Lys	Asp	Phe	Phe	Gly	Glu	Gly	Ser	Val	Arg	Glu	Ala	Thr	Ala	Lys	Leu
1															
														15	
Lys	Lys	Arg	Glu	Leu											
														20	

<210> 190

<211> 62

<212> PRT

<213> *Homo sapiens*

<400> 190

Lys	Lys	Arg	Pro	Glu	Asp	Phe	Lys	Phe	Gly	Lys	Ile	Leu	Gly	Glu	Gly
1															
														15	
Ser	Phe	Ser	Thr	Val	Val	Leu	Ala	Arg	Glu	Leu	Ala	Thr	Ser	Arg	Glu
														30	
Tyr	Ala	Ile	Lys	Ile	Leu	Glu	Lys	Arg	His	Ile	Ile	Lys	Glu	Asn	Lys
														45	
Val	Pro	Tyr	Val	Thr	Arg	Glu	Arg	Asp	Val	Met	Ser	Arg	Leu		
														60	
50															

<210> 191

<211> 90

<212> PRT

<213> *Caenorhabditis elegans*

<400> 191

His	Pro	Phe	Val	Thr	Gln	Leu	Tyr	Thr	His	Phe	His	Asp	Gln	Ala	Arg
1															
														15	
Ile	Tyr	Phe	Val	Ile	Gly	Leu	Val	Glu	Asn	Gly	Asp	Leu	Gly	Glu	Ser
														30	
Leu	Cys	His	Phe	Gly	Ser	Phe	Asp	Met	Leu	Thr	Ser	Lys	Phe	Phe	Ala
														45	
35															

Ser Glu Ile Leu Thr Gly Leu Gln Phe Leu His Asp Asn Lys Ile Val
50 55 60
His Arg Asp Met Lys Pro Asp Asn Val Leu Ile Gln Lys Asp Gly His
65 70 75 80
Ile Leu Ile Thr Asp Phe Gly Ser Ala Gln
85 90

<210> 192
<211> 39
<212> PRT
<213> *Caenorhabditis elegans*

<400> 192
His Pro Phe Leu Tyr Phe Asp Tyr Phe Asn Gly Leu Gly Ser Phe Asp
1 5 10 15
Phe Glu Ile Leu Leu His Ile His Arg Asp Lys Pro Asn Leu Asp His
20 25 30
Ile Ile Thr Asp Phe Gly Ala
35

<210> 193
<211> 90
<212> PRT
<213> *Homo sapiens*

<400> 193
His Pro Phe Phe Val Lys Leu Tyr Phe Thr Phe Gln Asp Asp Glu Lys
1 5 10 15
Leu Tyr Phe Gly Leu Ser Tyr Ala Lys Asn Gly Glu Leu Leu Lys Tyr
20 25 30
Ile Arg Lys Ile Gly Ser Phe Asp Glu Thr Cys Thr Arg Phe Tyr Thr
35 40 45
Ala Glu Ile Val Ser Ala Leu Glu Tyr Leu His Gly Lys Gly Ile Ile
50 55 60
His Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu Asn Glu Asp Met His
65 70 75 80
Ile Gln Ile Thr Asp Phe Gly Thr Ala Lys
85 90

<210> 194
<211> 98
<212> PRT
<213> *Caenorhabditis elegans*

<400> 194
Glu Glu Asn Thr Ala Arg Arg Thr Thr Phe Val Gly Thr Ala Leu Tyr
1 5 10 15
Val Ser Pro Glu Met Leu Ala Asp Gly Asp Val Gly Pro Gln Thr Asp
20 25 30
Ile Trp Gly Leu Gly Cys Ile Leu Phe Gln Cys Leu Ala Gly Gln Pro
35 40 45
Pro Phe Arg Ala Val Asn Gln Tyr His Leu Leu Lys Arg Ile Gln Glu
50 55 60
Leu Asp Phe Ser Phe Pro Glu Gly Phe Pro Glu Glu Ala Ser Glu Ile
65 70 75 80
Ile Ala Lys Ile Leu Val Arg Asp Pro Ser Thr Arg Ile Thr Ser Gln

85 90 95
Glu Leu

<210> 195
<211> 43
<212> PRT
<213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 195
Glu Ala Arg Phe Val Gly Thr Ala Tyr Val Ser Pro Glu Leu Asp Trp
1 5 10 15
Leu Gly Cys Ile Gln Ala Gly Pro Pro Phe Arg Ala Asn Tyr Ile Leu
20 25 30
Phe Pro Glu Phe Ala Lys Leu Val Asp Arg Glu
35 40

<210> 196
<211> 98
<212> PRT
<213> *Homo sapiens*

<400> 196
Glu Ser Lys Gln Ala Arg Ala Asn Ser Phe Val Gly Thr Ala Gln Tyr
1 5 10 15
Val Ser Pro Glu Leu Leu Thr Glu Lys Ser Ala Cys Lys Ser Ser Asp
20 25 30
Leu Trp Ala Leu Gly Cys Ile Ile Tyr Gln Leu Val Ala Gly Leu Pro
35 40 45
Pro Phe Arg Ala Gly Asn Glu Tyr Leu Ile Phe Gln Lys Ile Ile Lys
50 55 60
Leu Glu Tyr Asp Phe Pro Glu Lys Phe Phe Pro Lys Ala Arg Asp Leu
65 70 75 80
Val Glu Lys Leu Leu Val Leu Asp Ala Thr Lys Arg Leu Gly Cys Glu
85 90 95

Glu Met

<210> 197
<211> 35
<212> PRT
<213> *Caenorhabditis elegans*

<400> 197
Leu Met Ala His Lys Phe Phe Glu Asn Val Asp Trp Val Asn Ile Ala
1 5 10 15
Asn Ile Lys Pro Pro Val Leu His Ala Tyr Ile Pro Ala Thr Phe Gly
20 25 30
Glu Pro Glu
35

<210> 198
<211> 17
<212> PRT
<213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 198
Leu Ala His Phe Phe Glu Val Trp Asn Pro Pro Leu Ala Tyr Pro Ala
1 5 10 15
Glu

<210> 199
<211> 35
<212> PRT
<213> Homo sapiens

<400> 199
Leu Lys Ala His Pro Phe Phe Glu Ser Val Thr Trp Glu Asn Leu His
1 5 10 15
Gln Gln Thr Pro Pro Lys Leu Thr Ala Tyr Leu Pro Ala Met Ser Glu
20 25 30
Asp Asp Glu
35

<210> 200
<211> 104
<212> PRT
<213> Caenorhabditis elegans

<400> 200
Leu Glu Glu Gln Arg Val Lys Asn Pro Phe His Ile Phe Thr Asn Asn
1 5 10 15
Ser Leu Ile Leu Lys Gln Gly Tyr Leu Glu Lys Lys Arg Gly Leu Phe
20 25 30
Ala Arg Arg Arg Met Phe Leu Leu Thr Glu Gly Pro His Leu Leu Tyr
35 40 45
Ile Asp Val Pro Asn Leu Val Leu Lys Gly Glu Val Pro Trp Thr Pro
50 55 60
Cys Met Gln Val Glu Leu Lys Asn Ser Gly Thr Phe Phe Ile His Thr
65 70 75 80
Pro Asn Arg Val Tyr Tyr Leu Phe Asp Leu Glu Lys Lys Ala Asp Glu
85 90 95
Trp Cys Lys Ala Ile Asn Asp Val
100

<210> 201
<211> 59
<212> PRT
<213> Caenorhabditis elegans or Homo sapiens

<400> 201
Leu Glu Gln Asn Pro His Phe Asn Leu Ile Leu Lys Gly Lys Gly Leu
1 5 10 15
Phe Ala Arg Arg Arg Leu Leu Thr Glu Gly Pro His Leu Tyr Asp Asn
20 25 30
Val Leu Lys Gly Glu Pro Trp Glu Lys Asn Thr Phe Phe His Thr Pro
35 40 45
Asn Arg Tyr Tyr Leu Asp Ala Trp Cys Ile Val
50 55

<210> 202
<211> 104
<212> PRT
<213> Homo sapiens

<400> 202
Leu Glu Lys Gln Ala Gly Gly Asn Pro Trp His Gln Phe Val Glu Asn
1 5 10 15
Asn Leu Ile Leu Lys Met Gly Pro Val Asp Lys Arg Lys Gly Leu Phe
20 25 30
Ala Arg Arg Arg Gln Leu Leu Leu Thr Glu Gly Pro His Leu Tyr Tyr
35 40 45
Val Asp Pro Val Asn Lys Val Leu Lys Gly Glu Ile Pro Trp Ser Gln
50 55 60
Glu Leu Arg Pro Glu Ala Lys Asn Phe Lys Thr Phe Phe Val His Thr
65 70 75 80
Pro Asn Arg Thr Tyr Tyr Leu Met Asp Pro Ser Gly Asn Ala His Lys
85 90 95
Trp Cys Arg Lys Ile Gln Glu Val
100

<210> 203
<211> 45
<212> PRT
<213> Homo sapiens

<400> 203
Lys Leu Glu Asn Leu Met Leu Asp Lys Asp Gly His Ile Lys Ile Thr
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Gly Ile Lys Asp Gly Ala Thr Met Lys
20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 204
<211> 36
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans

<400> 204
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Asp Phe
1 5 10 15
Gly Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu
20 25 30
Ala Pro Glu Val
35

<210> 205
<211> 45
<212> PRT
<213> Caenorhabditis elegans

<400> 205
Lys Leu Glu Asn Leu Leu Asp Lys Asp Gly His Ile Lys Ile Ala
1 5 10 15
Asp Phe Gly Leu Cys Lys Glu Glu Ile Ser Phe Gly Asp Lys Thr Ser

20 25 30
Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val
35 40 45

<210> 206
<211> 62
<212> PRT
<213> *Caenorhabditis elegans*

<400> 206
Leu Cys Lys Glu Glu Ile Lys Tyr Gly Asp Lys Thr Ser Thr Phe Cys
1 5 10 15
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Ile Glu Asp Ile Asp Tyr
20 25 30
Asp Arg Ser Val Asp Trp Trp Gly Val Gly Val Val Met Tyr Glu Met
35 40 45
Met Cys Gly Arg Leu Pro Phe Ser Ala Lys Glu Asn Gly Lys
50 55 60

<210> 207
<211> 43
<212> PRT
<213> *Caenorhabditis elegans* or *Mus musculus*

<400> 207
Leu Cys Lys Glu Ile Gly Thr Phe Cys Gly Thr Pro Glu Tyr Leu Ala
1 5 10 15
Pro Glu Val Glu Asp Asp Tyr Arg Val Asp Trp Trp Gly Gly Val Val
20 25 30
Met Tyr Glu Met Met Cys Gly Arg Leu Pro Phe
35 40

<210> 208
<211> 492
<212> PRT
<213> *Caenorhabditis elegans*

<400> 208
Met Gly Val Asn Asp His Asp Val Ser Val Pro Leu Gln Glu Val Gln
1 5 10 15
Ser Arg Thr Val Glu Gly Lys Leu Thr Lys Cys Leu Ala Phe Ser Ala
20 25 30
Phe Val Ile Thr Leu Ala Ser Phe Gln Phe Gly Tyr His Ile Gly Cys
35 40 45
Val Asn Ala Pro Gly Gly Leu Ile Thr Glu Trp Ile Ile Gly Ser His
50 55 60
Lys Asp Leu Phe Asp Lys Glu Leu Ser Arg Glu Asn Ala Asp Leu Ala
65 70 75 80
Trp Ser Val Ala Val Ser Val Phe Ala Val Gly Gly Met Ile Gly Gly
85 90 95
Leu Ser Ser Gly Trp Leu Ala Asp Lys Val Gly Arg Arg Gly Ala Leu
100 105 110
Phe Tyr Asn Asn Leu Leu Ala Leu Ala Ala Ala Leu Met Gly Leu
115 120 125
Ala Lys Ser Val Gly Ala Tyr Pro Met Val Ile Leu Gly Arg Leu Ile
130 135 140

Ile	Gly	Leu	Asn	Cys	Gly	Phe	Ser	Ser	Ala	Leu	Val	Pro	Met	Phe	Leu
145					150					155					160
Thr	Glu	Ile	Ser	Pro	Asn	Asn	Leu	Arg	Gly	Met	Leu	Gly	Ser	Leu	His
					165				170						175
Gln	Leu	Leu	Val	Thr	Ile	Ala	Ile	Leu	Val	Ser	Gln	Ile	Phe	Gly	Leu
					180				185						190
Pro	His	Leu	Leu	Gly	Thr	Gly	Asp	Arg	Trp	Pro	Leu	Ile	Phe	Ala	Phe
					195			200							205
Thr	Val	Val	Pro	Ala	Val	Leu	Gln	Leu	Ala	Leu	Leu	Met	Leu	Cys	Pro
					210			215							220
Glu	Ser	Pro	Lys	Tyr	Thr	Met	Ala	Val	Arg	Gly	Gln	Arg	Asn	Glu	Ala
					225			230			235				240
Glu	Ser	Ala	Leu	Lys	Lys	Leu	Arg	Asp	Thr	Glu	Asp	Val	Ser	Thr	Glu
					245				250						255
Ile	Glu	Ala	Met	Gln	Glu	Glu	Ala	Thr	Ala	Ala	Gly	Val	Gln	Glu	Lys
					260			265							270
Pro	Lys	Met	Gly	Asp	Met	Phe	Lys	Gly	Ala	Leu	Leu	Trp	Pro	Met	Ser
					275			280							285
Ile	Ala	Ile	Met	Met	Met	Leu	Ala	Gln	Gln	Leu	Ser	Gly	Ile	Asn	Val
					290			295			300				
Ala	Met	Phe	Tyr	Ser	Thr	Val	Ile	Phe	Arg	Gly	Ala	Gly	Leu	Thr	Gly
					305			310			315				320
Asn	Glu	Pro	Phe	Tyr	Ala	Thr	Ile	Gly	Met	Gly	Ala	Val	Asn	Val	Ile
					325				330						335
Met	Thr	Leu	Ile	Ser	Val	Trp	Leu	Val	Asp	His	Pro	Lys	Phe	Gly	Arg
					340			345							350
Arg	Ser	Leu	Leu	Leu	Ala	Gly	Leu	Thr	Gly	Met	Phe	Val	Ser	Thr	Leu
					355			360							365
Leu	Leu	Val	Gly	Ala	Leu	Thr	Ile	Gln	Asn	Ser	Gly	Gly	Asp	Lys	Trp
					370			375			380				
Ala	Ser	Tyr	Ser	Ala	Ile	Gly	Phe	Val	Leu	Leu	Phe	Val	Ile	Ser	Phe
					385			390			395				400
Ala	Thr	Gly	Pro	Gly	Ala	Ile	Pro	Trp	Phe	Phe	Val	Ser	Glu	Ile	Phe
					405				410						415
Asp	Ser	Ser	Ala	Arg	Gly	Asn	Ala	Asn	Ser	Ile	Ala	Val	Met	Val	Asn
					420			425							430
Trp	Ala	Ala	Asn	Leu	Leu	Val	Gly	Leu	Thr	Phe	Leu	Pro	Ile	Asn	Asn
					435			440							445
Leu	Met	Gln	Gln	Tyr	Ser	Phe	Phe	Ile	Phe	Ser	Gly	Phe	Leu	Ala	Phe
					450			455			460				
Phe	Ile	Phe	Tyr	Thr	Trp	Lys	Phe	Val	Pro	Glu	Thr	Lys	Gly	Lys	Ser
					465			470			475				480
Ile	Glu	Gln	Ile	Gln	Ala	Glu	Phe	Glu	Lys	Arg	Lys				
					485				490						

<210> 209

<211> 22

<212> PRT

<213> *Caenorhabditis elegans*

<400> 209

Arg Asn Glu Ala Glu Ser Ala Leu Lys Lys Leu Arg Asp Thr Glu Asp

1 5 10 15

Val Ser Thr Glu Ile Glu

20

<210> 210

<211> 28
<212> DNA
<213> *Caenorhabditis elegans*

<400> 210
tctcggtt tgccgtcgga tgtctgcc

28

<210> 211
<211> 223
<212> PRT
<213> *Ascoris suum*

<400> 211
Ala Lys Asn Asn Gly Glu Phe Val Arg Cys Val His Ser Val Gly Gln
1 5 10 15
Pro Lys Pro Val Ala Thr Lys Val Ile Asn His Trp Pro Cys Asn Pro
20 25 30
Glu Lys Thr Ile Ile Ala His Arg Pro Ala Glu Arg Glu Ile Trp Ser
35 40 45
Phe Gly Ser Gly Tyr Gly Gly Asn Ser Leu Leu Gly Lys Lys Cys Phe
50 55 60
Ala Leu Arg Ile Ala Met Asn Ile Gly Tyr Asp Glu Gly Trp Met Ala
65 70 75 80
Glu His Met Leu Ile Met Gly Val Thr Ser Pro Lys Gly Glu Glu Arg
85 90 95
Phe Val Ala Ala Ala Phe Pro Ser Ala Cys Gly Lys Thr Asn Leu Ala
100 105 110
Met Leu Glu Pro Thr Ile Pro Gly Trp Lys Val Arg Val Ile Gly Asp
115 120 125
Asp Ile Ala Trp Met Lys Phe Gly Ala Asp Gly Arg Leu Tyr Ala Ile
130 135 140
Asn Pro Glu Tyr Gly Phe Phe Gly Val Ala Pro Gly Thr Ser His Lys
145 150 155 160
Thr Asn Pro Met Ala Met Ala Ser Phe Gln Glu Asn Thr Ile Phe Thr
165 170 175
Asn Val Ala Glu Thr Ala Asp Gly Glu Tyr Phe Trp Glu Gly Leu Glu
180 185 190
His Glu Val Lys Asn Pro Lys Val Asp Met Ile Asn Trp Leu Gly Glu
195 200 205
Pro Trp His Ile Gly Asp Glu Ser Lys Ala Ala His Pro Asn Ser
210 215 220

<210> 212
<211> 176
<212> PRT
<213> *Caenorhabditis elegans* or *Ascoris suum*

<400> 212
Ala Asn Phe Val Arg Cys His Ser Val Gly Pro Pro Val Val Ile Asn
1 5 10 15
His Trp Pro Cys Asn Pro Glu Ile Ala His Arg Pro Glu Arg Glu Ile
20 25 30
Trp Ser Phe Gly Ser Gly Tyr Gly Gly Asn Ser Leu Leu Gly Lys Lys
35 40 45
Cys Phe Ala Leu Arg Ile Ala Asn Ile Asp Glu Gly Trp Met Ala Glu
50 55 60
His Met Leu Ile Met Gly Val Thr Pro Gly Glu Phe Ala Ala Ala Phe
65 70 75 80

Pro	Ser	Ala	Cys	Gly	Lys	Thr	Asn	Leu	Ala	Met	Leu	Glu	Pro	Thr	Pro
85								90						95	
Gly	Trp	Lys	Val	Arg	Gly	Asp	Asp	Ile	Ala	Trp	Met	Lys	Phe	Gly	Asp
								105						110	
100															
Gly	Arg	Leu	Tyr	Ala	Ile	Asn	Pro	Glu	Gly	Phe	Phe	Gly	Val	Ala	Pro
								120					125		
115															
Gly	Thr	Ser	Lys	Thr	Asn	Pro	Met	Ala	Ala	Phe	Gln	Asn	Ile	Phe	Thr
								135			140				
130															
Asn	Val	Ala	Glu	Thr	Ala	Gly	Glu	Tyr	Phe	Trp	Glu	Gly	Leu	Glu	Glu
								150			155			160	
145															
Val	Asp	Trp	Leu	Gly	Glu	Trp	His	Ile	Gly	Ala	Ala	His	Pro	Asn	Ser
								165			170			175	

<210> 213
<211> 223
<212> PRT
<213> *Caenorhabditis elegans*

<400>	213														
Ala	Leu	Gly	Asn	Gln	Asp	Phe	Val	Arg	Cys	Ile	His	Ser	Val	Gly	Leu
1				5					10					15	
Pro	Arg	Pro	Val	Lys	Gln	Arg	Val	Ile	Asn	His	Trp	Pro	Cys	Asn	Pro
				20				25					30		
Glu	Arg	Val	Leu	Ile	Ala	His	Arg	Pro	Pro	Glu	Arg	Glu	Ile	Trp	Ser
				35			40				45				
Phe	Gly	Ser	Gly	Tyr	Gly	Gly	Asn	Ser	Leu	Leu	Gly	Lys	Lys	Cys	Phe
				50			55				60				
Ala	Leu	Arg	Ile	Ala	Ser	Asn	Ile	Ala	Lys	Asp	Glu	Gly	Trp	Met	Ala
				65			70			75			80		
Glu	His	Met	Leu	Ile	Met	Gly	Val	Thr	Arg	Pro	Cys	Gly	Arg	Glu	His
					85			90				95			
Phe	Ile	Ala	Ala	Ala	Phe	Pro	Ser	Ala	Cys	Gly	Lys	Thr	Asn	Leu	Ala
				100				105				110			
Met	Leu	Glu	Pro	Thr	Leu	Pro	Gly	Trp	Lys	Val	Arg	Cys	Val	Gly	Asp
				115				120			125				
Asp	Ile	Ala	Trp	Met	Lys	Phe	Gly	Glu	Asp	Gly	Arg	Leu	Tyr	Ala	Ile
				130			135			140					
Asn	Pro	Glu	Ala	Gly	Phe	Phe	Gly	Val	Ala	Pro	Gly	Thr	Ser	Asn	Lys
				145			150			155			160		
Thr	Asn	Pro	Met	Ala	Val	Ala	Thr	Phe	Gln	Lys	Asn	Ser	Ile	Phe	Thr
					165				170			175			
Asn	Val	Ala	Glu	Thr	Ala	Asn	Gly	Glu	Tyr	Phe	Trp	Glu	Gly	Leu	Glu
				180				185				190			
Asp	Glu	Ile	Ala	Asp	Lys	Asn	Val	Asp	Ile	Thr	Thr	Trp	Leu	Gly	Glu
				195				200			205				
Lys	Trp	His	Ile	Gly	Glu	Pro	Gly	Val	Ala	Ala	His	Pro	Asn	Ser	
				210				215			220				

<210> 214
<211> 173
<212> PRT
<213> *Ascoris suum*

<400>	214														
Lys	Gly	Asp	Phe	Val	Ser	Leu	Pro	Lys	His	Val	Gln	Arg	Phe	Val	Ala
1				5					10			15			
Glu	Lys	Ala	Glu	Leu	Met	Lys	Pro	Ser	Ala	Ile	Phe	Ile	Cys	Asp	Gly

20	25	30	
Ser Gln Asn Glu Ala Asp Glu	Leu Ile Ala Arg Cys Val	Glu Arg Gly	
35	40	45	
Val Leu Val Pro Leu Lys Ala	Tyr Lys Asn Asn	Tyr Leu Cys Arg Thr	
50	55	60	
Asp Pro Arg Asp Val Ala Arg Val	Glu Ser Lys Thr Trp Met	Ile Thr	
65	70	75	80
Pro Glu Lys Tyr Asp Ser Val	Cys His Thr Pro	Glu Gly Val Lys Pro	
85	90	95	
Met Met Gly Gln Trp Met Ser Pro	Asp Glu Phe Gly Lys	Glu Leu Asp	
100	105	110	
Asp Arg Phe Pro Gly Cys Met	Ala Gly Arg Thr Met	Tyr Val Ile Pro	
115	120	125	
Tyr Ser Met Gly Pro Val Gly	Pro Leu Ser Lys	Ile Gly Ile Glu	
130	135	140	
Leu Thr Asp Ser Asp Tyr Val	Val Leu Cys Met	Arg Ile Met Thr Arg	
145	150	155	160
Met Gly Glu Pro Val Leu Lys	Ala Leu Ala Lys	Asn Asn	
165	170		

<210> 215

<211> 120

<212> PRT

<213> *Caenorhabditis elegans* or *Ascoris suum*

<400> 215

Gly Asp Phe Leu Pro Val Gln Arg Phe	Ala Glu Lys Ala Glu	Leu Met	
1	5	10	15
Pro Ile Phe Ile Cys Asp Gly Ser	Gln Glu Ala Asp Glu	Leu Ile Glu	
20	25	30	
Arg Gly Leu Leu Ala Tyr Asn Asn	Tyr Cys Arg Thr	Asp Pro Asp Val	
35	40	45	
Ala Arg Val Glu Ser Lys Thr Trp	Met Thr Lys Tyr	Asp Val His Thr	
50	55	60	
Glu Gly Val Pro Met Gly Trp Pro	Glu Leu Asp Arg Phe	Pro Gly Cys	
65	70	75	80
Met Ala Gly Arg Met Tyr Val Ile	Pro Ser Met Gly Pro	Val Gly Gly	
85	90	95	
Pro Leu Ser Lys Ile Gly Ile Leu	Thr Asp Ser Tyr Val	Val Leu Met	
100	105	110	
Arg Ile Met Thr Arg Val Ala Leu			
115	120		

<210> 216

<211> 173

<212> PRT

<213> *Caenorhabditis elegans*

<400> 216

Gln Gly Asp Phe His Leu Leu Pro Ala Lys Val Gln Arg Phe	Ile Ala		
1	5	10	15
Glu Lys Ala Glu Leu Met Arg Pro Arg Gly	Ile Phe Ile Cys Asp Gly		
20	25	30	
Ser Gln His Glu Ala Asp Glu Leu Ile Asp Lys	Leu Ile Glu Arg Gly		
35	40	45	
Met Leu Ser Lys Leu Glu Ala Tyr Glu Asn Asn	Tyr Ile Cys Arg Thr		
50	55	60	

Asp	Pro	Lys	Asp	Val	Ala	Arg	Val	Glu	Ser	Lys	Thr	Trp	Met	Val	Thr
65				70					75				80		
Lys	Asn	Lys	Tyr	Asp	Thr	Val	Thr	His	Thr	Lys	Glu	Gly	Val	Glu	Pro
					85				90				95		
Ile	Met	Gly	His	Trp	Leu	Ala	Pro	Glu	Asp	Leu	Ala	Thr	Glu	Leu	Asp
					100			105				110			
Ser	Arg	Phe	Pro	Gly	Cys	Met	Ala	Gly	Arg	Ile	Met	Tyr	Val	Ile	Pro
					115			120			125				
Phe	Ser	Met	Gly	Pro	Val	Gly	Gly	Pro	Leu	Ser	Lys	Ile	Gly	Ile	Gln
					130			135			140				
Leu	Thr	Asp	Ser	Asn	Tyr	Val	Val	Leu	Ser	Met	Arg	Ile	Met	Thr	Arg
145					150				155			160			
Val	Asn	Asn	Asp	Val	Trp	Asp	Ala	Leu	Gly	Asn	Gln	Asp			
					165				170						

<210> 217

<211> 107

<212> PRT

<213> Ascoris suum

<400> 217

Arg	Phe	Thr	Ala	Pro	Ala	Gly	Gln	Cys	Pro	Ile	Ile	His	Pro	Asp	Trp
1					5				10			15			
Glu	Lys	Pro	Glu	Gly	Val	Pro	Ile	Asp	Ala	Ile	Ile	Phe	Gly	Gly	Arg
					20			25			30				
Arg	Pro	Glu	Gly	Val	Pro	Leu	Val	Phe	Glu	Ser	Arg	Ser	Trp	Val	His
					35			40			45				
Gly	Ile	Phe	Val	Gly	Ala	Cys	Val	Lys	Ser	Glu	Ala	Thr	Ala	Ala	Ala
					50			55			60				
Glu	His	Thr	Gly	Lys	Gln	Val	Met	His	Asp	Pro	Met	Ala	Met	Arg	Pro
65					70				75			80			
Phe	Met	Gly	Tyr	Asn	Phe	Gly	Arg	Tyr	Met	Arg	His	Trp	Met	Lys	Leu
					85			90			95				
Gly	Gln	Pro	Pro	His	Lys	Val	Pro	Lys	Ile	Phe					
					100				105						

<210> 218

<211> 77

<212> PRT

<213> Caenorhabditis elegans or Ascoris suum

<400> 218

Arg	Phe	Ala	Pro	Ala	Gln	Cys	Pro	Ile	Ile	His	Pro	Asp	Trp	Glu	Pro
1					5			10			15				
Gly	Val	Pro	Ile	Ala	Ile	Ile	Phe	Gly	Gly	Arg	Arg	Pro	Gly	Val	Pro
					20			25			30				
Leu	Glu	Ser	Trp	His	Gly	Phe	Gly	Cys	Lys	Ser	Glu	Ala	Thr	Ala	Ala
					35			40			45				
Ala	Glu	Thr	Gly	Lys	Val	Met	His	Asp	Pro	Met	Ala	Met	Arg	Pro	Phe
					50			55			60				
Met	Gly	Tyr	Asn	Phe	Gly	Tyr	His	Trp	Leu	Lys	Val	Phe			
65					70				75						

<210> 219

<211> 107

<212> PRT

<213> *Caenorhabditis elegans*

<400> 219

Arg	Phe	Ala	Ala	Pro	Ala	Asn	Gln	Cys	Pro	Ile	Ile	His	Pro	Asp	Trp
1															
															15
Glu	Ser	Pro	Gln	Gly	Val	Pro	Ile	Glu	Ala	Ile	Ile	Phe	Gly	Gly	Arg
															30
Arg	Pro	Gln	Gly	Val	Pro	Leu	Ile	Tyr	Glu	Thr	Asn	Ser	Trp	Glu	His
															45
Gly	Val	Phe	Thr	Gly	Ser	Cys	Leu	Lys	Ser	Glu	Ala	Thr	Ala	Ala	Ala
															60
Glu	Phe	Thr	Gly	Lys	Thr	Val	Met	His	Asp	Pro	Met	Ala	Met	Arg	Pro
															80
Phe	Met	Gly	Tyr	Asn	Phe	Gly	Lys	Tyr	Leu	Gln	His	Trp	Leu	Asp	Leu
															95
Lys	Thr	Asp	Ser	Arg	Lys	Val	Ile	Asp	Phe	Phe					
100															
105															

<210> 220

<211> 116

<212> PRT

<213> *Ascoris suum*

<400> 220

Val	Pro	Lys	Ile	Phe	His	Val	Asn	Trp	Phe	Arg	Gln	Ser	Ala	Asp	His
1															
															15
Lys	Phe	Leu	Trp	Pro	Gly	Tyr	Gly	Asp	Asn	Ile	Arg	Val	Ile	Asp	Trp
															30
Ile	Leu	Arg	Arg	Cys	Ser	Gly	Asp	Ala	Thr	Ile	Ala	Glu	Glu	Thr	Pro
															45
Ile	Gly	Phe	Ile	Pro	Lys	Lys	Gly	Thr	Ile	Asn	Leu	Glu	Gly	Leu	Pro
															60
Asn	Val	Asn	Trp	Asp	Glu	Leu	Met	Ser	Ile	Pro	Lys	Ser	Tyr	Trp	Leu
															80
Glu	Asp	Met	Val	Glu	Thr	Lys	Thr	Phe	Phe	Glu	Asn	Gln	Val	Gly	Ser
															95
Asp	Leu	Pro	Pro	Glu	Ile	Ala	Lys	Glu	Leu	Glu	Ala	Gln	Thr	Glu	Arg
															110
Ile	Lys	Ala	Leu												
115															

<210> 221

<211> 68

<212> PRT

<213> *Caenorhabditis elegans* or *Ascoris suum*

<400> 221

Pro	Lys	Ile	His	Val	Asn	Trp	Phe	Arg	Lys	Phe	Leu	Trp	Pro	Gly	Gly
1															
															15
Asp	Asn	Ile	Arg	Val	Ile	Asp	Trp	Ile	Arg	Arg	Gly	Ile	Glu	Thr	Pro
															30
Ile	Gly	Pro	Lys	Gly	Ile	Asn	Leu	Glu	Gly	Leu	Val	Asn	Trp	Asp	Glu
															45
Leu	Met	Ser	Pro	Tyr	Trp	Asp	Glu	Phe	Gln	Val	Gly	Asp	Leu	Pro	Glu
															60
Ala	Gln	Arg	Leu												
65															

<210> 222
<211> 116
<212> PRT
<213> *Caenorhabditis elegans*

<400> 222
Met Pro Lys Ile Tyr His Val Asn Trp Phe Arg Lys Asp Ser Asn Asn
1 5 10 15
Lys Phe Leu Trp Pro Gly Phe Gly Asp Asn Ile Arg Val Ile Asp Trp
20 25 30
Ile Ile Arg Arg Leu Asp Gly Glu Gln Glu Ile Gly Val Glu Thr Pro
35 40 45
Ile Gly Thr Val Pro Ala Lys Gly Ser Ile Asn Leu Glu Gly Leu Gly
50 55 60
Glu Val Asn Trp Asp Glu Leu Met Ser Val Pro Ala Asp Tyr Trp Lys
65 70 75 80
Gln Asp Ala Gln Glu Ile Arg Lys Phe Leu Asp Glu Gln Val Gly Glu
85 90 95
Asp Leu Pro Glu Pro Val Arg Ala Glu Met Asp Ala Gln Glu Lys Arg
100 105 110
Val Gln Thr Leu
115

<210> 223
<211> 36
<212> PRT
<213> *Ascoris suum*

<400> 223
Ser Leu Ser His Phe Lys Asp Asp Asp Phe Ala Val Val Ser Glu Val
1 5 10 15
Val Thr His Lys Gln Asn His Ile Pro Val Ile Lys Gly Asp Phe Val
20 25 30
Ser Leu Pro Lys
35

<210> 224
<211> 15
<212> PRT
<213> *Caenorhabditis elegans* or *Ascoris suum*

<400> 224
Ser Leu Asp Phe Val Val Glu Val Val His Pro Lys Phe Ser Lys
1 5 10 15

<210> 225
<211> 36
<212> PRT
<213> *Caenorhabditis elegans*

<400> 225
Ser Leu Arg Gln Ile Ser Glu Asp Ala Phe Tyr Val Val Asn Glu Val
1 5 10 15
Val Met Lys Arg Leu Gly His Val Pro Ile Leu Lys Val Ile Phe Glu
20 25 30

Ser Ser Glu Lys
35

<210> 226
<211> 25
<212> PRT
<213> Ascoris suum

<400> 226
Gly Cys Met Ala Gly Arg Thr Met Tyr Val Ile Pro Tyr Ser Met Gly
1 5 10 15
Pro Val Gly Gly Pro Leu Ser Lys Ile
20 25

<210> 227
<211> 9
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum

<400> 227
Gly Cys Arg Val Pro Ser Pro Leu Lys
1 5

<210> 228
<211> 25
<212> PRT
<213> Caenorhabditis elegans

<400> 228
Gly Cys Ser Gly Arg Arg Val Leu Cys Val Cys Pro Cys Ser His Ser
1 5 10 15
Ser Ser Ala Leu Pro Leu Gln Lys Val
20 25

<210> 229
<211> 16
<212> PRT
<213> Ascoris suum

<400> 229
Leu Pro Asn Val Asn Trp Asp Glu Leu Met Ser Ile Pro Lys Ser Tyr
1 5 10 15

<210> 230
<211> 7
<212> PRT
<213> Caenorhabditis elegans or Ascoris suum

<400> 230
Leu Asn Trp Ser Pro Ser Tyr
1 5

<210> 231

<211> 16

<212> PRT

<213> *Caenorhabditis elegans*

<400> 231

Leu Glu Ser Phe Asn Trp Phe Ser Phe Val Ser Cys Pro Asp Ser Tyr
1 5 10 15

<210> 232

<211> 14

<212> PRT

<213> *Ascoris suum*

<400> 232

Ser Val Cys His Thr Pro Glu Gly Val Lys Pro Met Met Gly
1 5 10

<210> 233

<211> 6

<212> PRT

<213> *Caenorhabditis elegans* or *Ascoris suum*

<400> 233

Val His Pro Pro Met Gly
1 5

<210> 234

<211> 14

<212> PRT

<213> *Caenorhabditis elegans*

<400> 234

Thr Val Met His Asp Pro Met Ala Met Arg Pro Phe Met Gly
1 5 10

<210> 235

<211> 197

<212> PRT

<213> *Homo sapiens*

<400> 235

Ser Gly Phe Phe Asp Tyr Gly Ser Phe Ser Glu Ile Met Gln Pro Trp
1 5 10 15

Ala Gln Thr Val Val Val Gly Arg Ala Arg Leu Gly Gly Ile Pro Val
20 25 30

Gly Val Val Ala Val Glu Thr Arg Thr Val Glu Leu Ser Val Pro Ala
35 40 45

Asp Pro Ala Asn Leu Asp Ser Glu Ala Lys Ile Ile Gln Gln Ala Gly
50 55 60

Gln Val Trp Phe Pro Asp Ser Ala Phe Lys Thr Tyr Gln Ala Ile Lys
65 70 75 80

Asp Phe Asn Arg Glu Gly Leu Pro Leu Met Val Phe Ala Asn Trp Arg
85 90 95

Gly Phe Ser Gly Gly Met Lys Asp Met Tyr Asp Gln Val Leu Lys Phe
100 105 110

Gly Ala Tyr Ile Val Asp Gly Leu Arg Glu Cys Ser Gln Pro Val Met
 115 120 125
 Val Tyr Ile Pro Pro Gln Ala Glu Leu Arg Gly Gly Ser Trp Val Val
 130 135 140
 Ile Asp Pro Thr Ile Asn Pro Arg His Met Glu Met Tyr Ala Asp Arg
 145 150 155 160
 Glu Ser Arg Gly Ser Val Leu Glu Pro Glu Gly Thr Val Glu Ile Lys
 165 170 175
 Phe Arg Lys Lys Asp Leu Val Lys Thr Met Arg Arg Val Asp Pro Val
 180 185 190
 Tyr Ile Arg Leu Ala
 195

<210> 236
 <211> 109
 <212> PRT
 <213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 236
 Gly Asp Ser Phe Glu Ile Trp Ala Val Gly Arg Ala Arg Leu Gly Ile
 1 5 10 15
 Pro Gly Val Val Glu Arg Val Pro Ala Asp Pro Ala Ser Gln Ala Gly
 20 25 30
 Gln Val Trp Pro Asp Ser Ala Phe Lys Thr Ala Ile Asp Asn Glu Leu
 35 40 45
 Pro Leu Met Ala Arg Gly Phe Ser Gly Gly Lys Asp Met Tyr Asp Val
 50 55 60
 Leu Lys Phe Gly Ala Ile Val Asp Leu Pro Val Val Tyr Ile Pro Glu
 65 70 75 80
 Leu Arg Gly Gly Trp Val Asp Ile Pro Ala Asp Ser Arg Gly Leu Glu
 85 90 95
 Pro Val Ile Lys Phe Arg Lys Met Arg Asp Pro Tyr Leu
 100 105

<210> 237
 <211> 197
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 237
 Thr Gly Ile Cys Asp Thr Met Ser Phe Asp Glu Ile Cys Gly Asp Trp
 1 5 10 15
 Ala Lys Ser Ile Val Ala Gly Arg Ala Arg Leu Cys Gly Ile Pro Ile
 20 25 30
 Gly Val Val Ser Ser Glu Phe Arg Asn Phe Ser Thr Ile Val Pro Ala
 35 40 45
 Asp Pro Ala Ile Asp Gly Ser Gln Val Gln Asn Thr Gln Arg Ala Gly
 50 55 60
 Gln Val Trp Tyr Pro Asp Ser Ala Phe Lys Thr Ala Glu Ala Ile Asn
 65 70 75 80
 Asp Leu Asn Lys Glu Asn Leu Pro Leu Met Ile Ile Ala Ser Leu Arg
 85 90 95
 Gly Phe Ser Gly Gly Gln Lys Asp Met Tyr Asp Met Val Leu Lys Phe
 100 105 110
 Gly Ala Gln Ile Val Asp Ala Leu Ala Val Tyr Asn Arg Pro Val Ile
 115 120 125
 Val Tyr Ile Pro Glu Ala Gly Glu Leu Arg Gly Gly Ala Trp Ala Val

130	135	140
Leu Asp Ser Lys Ile Arg Pro Glu Phe Ile His	Leu Val Ala Asp Glu	
145	150	155 160
Lys Ser Arg Gly Gly Ile Leu Glu Pro Asn Ala Val Val Gly Ile Lys		
165	170	175
Phe Arg Lys Pro Met Met Glu Met Met Lys Arg Ser Asp Pro Thr		
180	185	190
Tyr Ser Lys Leu Ser		
195		

<210> 238
<211> 124
<212> PRT
<213> Homo sapiens

<220>
<221> VARIANT
<222> (1)...(124)
<223> Xaa = Any Amino Acid

<400> 238
Val Gly Tyr Pro Val Met Ile Lys Ala Ser Glu Gly Gly Gly Lys
1 5 10 15
Gly Ile Arg Lys Val Asn Asn Ala Asp Asp Phe Pro Asn Leu Phe Arg
20 25 30
Gln Val Gln Ala Glu Val Pro Gly Ser Pro Ile Phe Val Met Arg Leu
35 40 45
Ala Lys Gln Ser Arg His Leu Glu Val Gln Ile Leu Ala Asp Gln Tyr
50 55 60
Gly Asn Ala Ile Ser Leu Phe Gly Arg Asp Cys Ser Val Gln Arg Arg
65 70 75 80
His Gln Lys Xaa
85 90 95
Val Phe Glu His Met Glu Gln Cys Ala Val Lys Leu Ala Lys Met Val
100 105 110
Gly Tyr Val Ser Ala Gly Thr Val Glu Tyr Leu Tyr
115 120

<210> 239
<211> 68
<212> PRT
<213> Homo sapiens or Caenorhabditis elegans

<400> 239
Gly Pro Met Ile Lys Ala Ser Glu Gly Gly Gly Lys Gly Ile Arg
1 5 10 15
Lys Asp Phe Phe Val Glu Val Gly Ser Pro Ile Phe Met Arg His Glu
20 25 30
Val Gln Leu Ala Asp Tyr Asn Ile Ser Arg Asp Cys Ser Gln Arg Arg
35 40 45
Gln Lys Met Ala Val Leu Ala Lys Val Gly Tyr Ser Ala Gly Thr Val
50 55 60
Glu Tyr Leu Tyr
65

<210> 240

<211> 124
<212> PRT
<213> *Caenorhabditis elegans*

<400> 240

Ile	Gly	Phe	Pro	Leu	Met	Ile	Lys	Ala	Ser	Glu	Gly	Gly	Gly	Lys	
1				5				10						15	
Gly	Ile	Arg	Lys	Cys	Thr	Lys	Val	Glu	Asp	Phe	Lys	Ser	Met	Phe	Glu
				20				25						30	
Glu	Val	Ala	Gln	Glu	Val	Gln	Gly	Ser	Pro	Ile	Phe	Leu	Met	Lys	Cys
				35			40					45			
Val	Asp	Gly	Ala	Arg	His	Ile	Glu	Val	Gln	Leu	Leu	Ala	Asp	Arg	Tyr
				50			55			60					
Glu	Asn	Val	Ile	Ser	Val	Tyr	Thr	Arg	Asp	Cys	Ser	Ile	Gln	Arg	Arg
				65		70		75					80		
Cys	Gln	Lys	Ile	Ile	Glu	Glu	Ala	Pro	Ala	Ile	Ile	Ala	Ser	Ser	His
				85			90					95			
Ile	Arg	Lys	Ser	Met	Gln	Glu	Asp	Ala	Val	Arg	Leu	Ala	Lys	Tyr	Val
				100			105			110					
Gly	Tyr	Glu	Ser	Ala	Gly	Thr	Val	Glu	Tyr	Leu	Tyr				
				115			120								

<210> 241
<211> 116
<212> PRT
<213> Rat

<400> 241

Lys	Glu	Glu	Gly	Leu	Gly	Ala	Glu	Asn	Leu	Arg	Gly	Ser	Gly	Met	Ile
1				5				10						15	
Ala	Gly	Glu	Ser	Ser	Leu	Ala	Tyr	Asp	Glu	Ile	Ile	Thr	Ile	Ser	Leu
				20				25				30			
Val	Thr	Cys	Arg	Ala	Ile	Gly	Ile	Gly	Ala	Tyr	Leu	Val	Arg	Leu	Gly
				35			40				45				
Gln	Arg	Thr	Ile	Gln	Val	Glu	Asn	Ser	His	Leu	Ile	Leu	Thr	Gly	Ala
				50		55			60						
Gly	Ala	Leu	Asn	Lys	Val	Leu	Gly	Arg	Glu	Val	Tyr	Thr	Ser	Asn	Asn
				65		70		75			80				
Gln	Leu	Gly	Gly	Ile	Gln	Ile	Met	His	Asn	Asn	Gly	Val	Thr	His	Cys
				85			90				95				
Thr	Val	Cys	Asp	Asp	Phe	Glu	Gly	Val	Phe	Thr	Val	Leu	His	Trp	Leu
				100			105			110					
Ser	Tyr	Met	Pro												
				115											

<210> 242
<211> 65
<212> PRT
<213> *Caenorhabditis elegans* or Rat

<400> 242

Lys	Glu	Gly	Glu	Asn	Leu	Gly	Ser	Gly	Ile	Ala	Gly	Glu	Ala	Tyr	Glu
1				5				10						15	
Thr	Val	Thr	Arg	Gly	Ile	Gly	Ala	Tyr	Arg	Leu	Arg	Gln	Ser	His	Leu
				20				25			30				
Ile	Leu	Thr	Gly	Ala	Leu	Asn	Leu	Gly	Val	Tyr	Thr	Ser	Asn	Asn	Gln
				35			40			45					

Leu Gly Gly Met Asn Gly Val Thr His Val Asp Glu Gly Val Trp Ser
50 55 60
Pro
65

<210> 243
<211> 116
<212> PRT
<213> *Caenorhabditis elegans*

<400> 243
Lys Asn Glu Lys Ile Gly Val Glu Asn Leu Gln Gly Ser Gly Leu Ile
1 5 10 15
Ala Gly Glu Thr Ala Arg Ala Tyr Ala Glu Val Pro Thr Tyr Cys Tyr
20 25 30
Val Thr Gly Arg Ser Val Gly Ile Gly Ala Tyr Thr Ala Arg Leu Ala
35 40 45
His Arg Ile Val Gln His Lys Gln Ser His Leu Ile Leu Thr Gly Tyr
50 55 60
Glu Ala Leu Asn Thr Leu Leu Gly Lys Lys Val Tyr Thr Ser Asn Asn
65 70 75 80
Gln Leu Gly Gly Pro Glu Val Met Phe Arg Asn Gly Val Thr His Ala
85 90 95
Val Val Asp Asn Asp Leu Glu Gly Ile Ala Lys Val Ile Arg Trp Met
100 105 110
Ser Phe Leu Pro
115

<210> 244
<211> 119
<212> PRT
<213> *Homo sapiens*

<400> 244
His Val Ile Ala Ala Arg Ile Thr Ser Glu Asn Pro Asp Glu Gly Phe
1 5 10 15
Lys Pro Ser Ser Gly Thr Val Gln Glu Leu Asn Phe Arg Ser Asn Lys
20 25 30
Asn Val Trp Gly Tyr Phe Ser Val Ala Ala Ala Gly Gly Leu His Glu
35 40 45
Phe Ala Asp Ser Gln Phe Gly His Cys Phe Ser Trp Gly Glu Asn Arg
50 55 60
Glu Glu Ala Ile Ser Asn Met Val Val Ala Leu Lys Glu Leu Ser Ile
65 70 75 80
Arg Gly Asp Phe Arg Thr Thr Val Glu Tyr Leu Ile Lys Leu Leu Glu
85 90 95
Thr Glu Ser Phe Gln Leu Asn Arg Ile Asp Thr Gly Trp Leu Asp Arg
100 105 110
Leu Ile Ala Glu Lys Val Gln
115

<210> 245
<211> 59
<212> PRT
<213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 245
His Ile Ala Ala Arg Ile Thr Glu Asn Pro Asp Phe Pro Ser Gly Val
1 5 10 15
Glu Asn Phe Ser Trp Tyr Phe Ser Val His Phe Ala Asp Ser Gln Phe
20 25 30
Gly His Phe Gly Arg Glu Ala Met Leu Lys Ile Arg Phe Thr Val Tyr
35 40 45
Leu Leu Phe Asn Thr Trp Leu Asp Ile Ala Lys
50 55

<210> 246
<211> 119
<212> PRT
<213> *Caenorhabditis elegans*

<400> 246
His Ala Ile Ala Ala Arg Ile Thr Cys Glu Asn Pro Asp Asp Ser Phe
1 5 10 15
Arg Pro Ser Thr Gly Lys Val Tyr Glu Ile Asn Phe Pro Ser Ser Gln
20 25 30
Asp Ala Trp Ala Tyr Phe Ser Val Gly Arg Gly Ser Ser Val His Gln
35 40 45
Phe Ala Asp Ser Gln Phe Gly His Ile Phe Thr Arg Gly Thr Ser Arg
50 55 60
Thr Glu Ala Met Asn Thr Met Cys Ser Thr Leu Lys His Met Thr Ile
65 70 75 80
Arg Ser Ser Phe Pro Thr Gln Val Asn Tyr Leu Val Asp Leu Met His
85 90 95
Asp Ala Asp Phe Ile Asn Asn Ala Phe Asn Thr Gln Trp Leu Asp Lys
100 105 110
Arg Ile Ala Met Lys Ile Lys
115

<210> 247
<211> 90
<212> PRT
<213> Rat

<400> 247
Pro Gly Gly Ala Asn Asn Asn Asn Tyr Ala Asn Val Glu Leu Ile Leu
1 5 10 15
Asp Ile Ala Lys Arg Ile Pro Val Gln Ala Val Trp Ala Gly Trp Gly
20 25 30
His Ala Ser Glu Asn Pro Lys Leu Pro Glu Leu Leu Lys Asn Gly
35 40 45
Ile Ala Phe Met Gly Pro Pro Ser Gln Ala Met Trp Ala Leu Gly Asp
50 55 60
Lys Ile Ala Ser Ser Ile Val Ala Gln Thr Ala Gly Ile Pro Thr Leu
65 70 75 80
Pro Trp Ser Gly Ser Gly Leu Arg Val Asp
85 90

<210> 248
<211> 55
<212> PRT
<213> *Caenorhabditis elegans* or Rat

<400> 248
Pro Gly Asn Asn Asn Ala Asn Val Ile Leu Ala Val Ala Val Trp Ala
1 5 10 15
Gly Trp Gly His Ala Ser Glu Asn Pro Leu Pro Leu Ile Ala Phe Gly
20 25 30
Pro Pro Ala Met Leu Gly Asp Lys Ile Ala Ser Ile Ala Gln Thr Gly
35 40 45
Pro Thr Trp Ser Gly Ser Gly
50 55

<210> 249
<211> 90
<212> PRT
<213> *Caenorhabditis elegans*

<400> 249
Pro Ser Gly Thr Asn Lys Asn Asn Phe Ala Asn Val Asp Glu Ile Leu
1 5 10 15
Lys His Ala Ile Lys Tyr Glu Val Asp Ala Val Trp Ala Gly Trp Gly
20 25 30
His Ala Ser Glu Asn Pro Asp Leu Pro Arg Arg Leu Asn Asp His Asn
35 40 45
Ile Ala Phe Ile Gly Pro Pro Ala Ser Ala Met Phe Ser Leu Gly Asp
50 55 60
Lys Ile Ala Ser Thr Ile Ile Ala Gln Thr Val Gly Val Pro Thr Val
65 70 75 80
Ala Trp Ser Gly Ser Gly Ile Thr Met Glu
85 90

<210> 250
<211> 67
<212> PRT
<213> *Caenorhabditis elegans*

<400> 250
Val Ile Lys Asn Leu Gly Tyr Met Val Asp Asn His Gly Phe Val Pro
1 5 10 15
Asn Gly Gly Arg Val Tyr Tyr Leu Thr Arg Ser Gln Pro Pro Leu Leu
20 25 30
Thr Pro Met Val Tyr Glu Tyr Tyr Met Ser Thr Gly Asp Leu Asp Phe
35 40 45
Val Met Glu Ile Leu Pro Thr Leu Asp Lys Glu Tyr Glu Phe Trp Ile
50 55 60
Lys Asn Arg
65

<210> 251
<211> 36
<212> PRT
<213> *Caenorhabditis elegans*

<400> 251
Ile Asn Gly Phe Val Pro Asn Gly Gly Arg Val Tyr Tyr Leu Arg Ser
1 5 10 15
Gln Pro Pro Pro Met Val Tyr Glu Tyr Tyr Thr Asp Val Pro Lys Glu
20 25 30

Tyr Phe Trp Arg
35

<210> 252
<211> 67
<212> PRT
<213> *Caenorhabditis elegans*

<400> 252
Met Ile Leu Asn Phe Ala His Ile Ile Glu Thr Tyr Gly Phe Val Pro
1 5 10 15
Asn Gly Gly Arg Val Tyr Tyr Leu Arg Arg Ser Gln Pro Pro Phe Phe
20 25 30
Ala Pro Met Val Tyr Glu Tyr Tyr Leu Ala Thr Gln Asp Ile Gln Leu
35 40 45
Val Ala Asp Leu Ile Pro Val Ile Glu Lys Glu Tyr Thr Phe Trp Ser
50 55 60
Glu Arg Arg
65

<210> 253
<211> 92
<212> PRT
<213> *Caenorhabditis elegans*

<400> 253
Met Asp Ser Ile Arg Thr Trp Ser Ile Ile Pro Ala Asp Leu Asn Ala
1 5 10 15
Phe Met Cys Ala Asn Ala Arg Ile Leu Ala Ser Leu Tyr Glu Ile Ala
20 25 30
Gly Asp Phe Lys Lys Val Lys Val Phe Glu Gln Arg Tyr Thr Trp Ala
35 40 45
Lys Arg Glu Met Arg Glu Leu His Trp Asn Glu Thr Asp Gly Ile Trp
50 55 60
Tyr Asp Tyr Asp Ile Glu Leu Lys Thr His Ser Asn Gln Tyr Tyr Val
65 70 75 80
Ser Asn Ala Val Pro Leu Tyr Ala Lys Cys Tyr Asp
85 90

<210> 254
<211> 32
<212> PRT
<213> *Caenorhabditis elegans*

<400> 254
Ile Thr Ile Pro Asp Leu Asn Ala Phe Cys Asn Ile Tyr Gly Lys Arg
1 5 10 15
Thr Trp Tyr Asp Tyr Thr His Ser Asn Ala Val Pro Leu Cys Tyr Asp
20 25 30

<210> 255
<211> 92
<212> PRT
<213> *Caenorhabditis elegans*

<400> 255
Ile Ser Thr Ile Glu Thr Thr Asn Ile Val Pro Val Asp Leu Asn Ala
1 5 10 15
Phe Leu Cys Tyr Asn Met Asn Ile Met Gln Leu Phe Tyr Lys Leu Thr
20 25 30
Gly Asn Pro Leu Lys His Leu Glu Trp Ser Ser Arg Phe Thr Asn Phe
35 40 45
Arg Glu Ala Phe Thr Lys Val Phe Tyr Val Pro Ala Arg Lys Gly Trp
50 55 60
Tyr Asp Tyr Asn Leu Arg Thr Leu Thr His Asn Thr Asp Phe Phe Ala
65 70 75 80
Ser Asn Ala Val Pro Leu Phe Ser Gln Cys Tyr Asp
85 90

<210> 256
<211> 102
<212> PRT
<213> *Caenorhabditis elegans*

<400> 256
Val His Asp Tyr Leu Glu Arg Gln Gly Leu Leu Lys Tyr Thr Lys Gly
1 5 10 15
Leu Pro Thr Ser Leu Ala Met Ser Ser Thr Gln Gln Trp Asp Lys Glu
20 25 30
Asn Ala Trp Pro Pro Met Ile His Met Val Ile Glu Gly Phe Arg Thr
35 40 45
Thr Gly Asp Ile Lys Leu Met Lys Val Ala Glu Lys Met Ala Thr Ser
50 55 60
Trp Leu Thr Gly Thr Tyr Gln Ser Phe Ile Arg Thr His Ala Met Phe
65 70 75 80
Glu Lys Tyr Asn Val Thr Pro His Thr Glu Glu Thr Ser Gly Gly Gly
85 90 95
Gly Gly Glu Tyr Glu Val
100

<210> 257
<211> 37
<212> PRT
<213> *Caenorhabditis elegans*

<400> 257
Val Gly Gly Pro Thr Ser Gln Gln Trp Asp Asn Trp Pro Met His Met
1 5 10 15
Ile Glu Gly Arg Leu Ala Ala Trp Leu Gln Phe Met Glu Lys Tyr Asn
20 25 30
Val Gly Gly Glu Val
35

<210> 258
<211> 102
<212> PRT
<213> *Caenorhabditis elegans*

<400> 258
Val Tyr Asn Glu Met Gln Asn Ser Gly Ala Phe Ser Ile Pro Gly Gly
1 5 10 15

Ile Pro Thr Ser Met Asn Glu Glu Thr Asn Gln Gln Trp Asp Phe Pro
20 25 30
Asn Gly Trp Ser Pro Met Asn His Met Ile Ile Glu Gly Leu Arg Lys
35 40 45
Ser Asn Asn Pro Ile Leu Gln Gln Lys Ala Phe Thr Leu Ala Glu Lys
50 55 60
Trp Leu Glu Thr Asn Met Gln Thr Phe Asn Val Ser Asp Glu Met Trp
65 70 75 80
Glu Lys Tyr Asn Val Lys Glu Pro Leu Gly Lys Leu Ala Thr Gly Gly
85 90 95
Glu Tyr Glu Val Gln Val
100

<210> 259

<211> 58

<212> PRT

<213> *Caenorhabditis elegans*

<400> 259

Tyr Gln Tyr Lys Ala Lys Leu Lys Val Pro Arg Pro Glu Ser Tyr Arg
1 5 10 15
Glu Asp Ser Glu Leu Ala Glu His Leu Gln Thr Glu Ala Glu Lys Ile
20 25 30
Gln Met Trp Ser Glu Ile Ala Ser Ala Ala Glu Thr Gly Trp Asp Phe
35 40 45
Ser Thr Arg Trp Phe Ser Gln Asn Gly Asp
50 55

<210> 260

<211> 29

<212> PRT

<213> *Caenorhabditis elegans*

<400> 260

Gln Tyr Pro Arg Pro Glu Ser Arg Glu Asp Ala Glu His Thr Lys Gln
1 5 10 15
Ser Ala Ala Glu Gly Trp Asp Phe Ser Arg Trp Phe Asp
20 25

<210> 261

<211> 58

<212> PRT

<213> *Caenorhabditis elegans*

<400> 261

Phe Gln Tyr Arg Thr Glu Ala Glu Thr Pro Arg Pro Glu Ser Phe Arg
1 5 10 15
Glu Asp Val Leu Ser Ala Glu His Phe Thr Thr Lys Asp Arg Lys Lys
20 25 30
Gln Phe Phe Lys Asp Leu Gly Ser Ala Ala Glu Ser Gly Trp Asp Phe
35 40 45
Ser Ser Arg Trp Phe Lys Asn His Lys Asp
50 55

<210> 262

<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 262
Gln Thr Gly Phe Gly Trp Thr Asn Gly Val Ile Leu Asp Leu Leu Asp
1 5 10 15
Lys Tyr Gly Asp Gln
20

<210> 263
<211> 13
<212> PRT
<213> *Caenorhabditis elegans*

<400> 263
Gln Gly Phe Gly Trp Thr Asn Gly Leu Asp Leu Tyr Asp
1 5 10

<210> 264
<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 264
Gln Ala Gly Phe Gly Trp Thr Asn Gly Ala Ala Leu Asp Leu Ile Phe
1 5 10 15
Thr Tyr Ser Asp Arg
20

<210> 265
<211> 24
<212> PRT
<213> *Caenorhabditis elegans*

<400> 265
Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe Ser Leu Ser Asn Ile Thr
1 5 10 15
Phe Val Val Phe Ile Leu Tyr Ile
20

<210> 266
<211> 10
<212> PRT
<213> *Caenorhabditis elegans*

<400> 266
Ser Ser Ser Phe Ser Val Phe Leu Tyr Ile
1 5 10

<210> 267
<211> 24
<212> PRT
<213> *Caenorhabditis elegans*

<400> 267
Thr Ser Ser Ser Ser Thr Phe Gly Tyr Ser Asn Ile Leu Thr Leu
1 5 10 15
Ile Thr Val Phe Val Leu Tyr Ile
20

<210> 268
<211> 7
<212> PRT
<213> *Caenorhabditis elegans*

<400> 268
Gly Gly Glu Tyr Glu Val Gln
1 5

<210> 269
<211> 7
<212> PRT
<213> *Caenorhabditis elegans*

<400> 269
Gly Gly Glu Tyr Glu Val Gln
1 5

<210> 270
<211> 7
<212> PRT
<213> *Caenorhabditis elegans*

<400> 270
Gly Gly Glu Tyr Glu Val Gln
1 5

<210> 271
<211> 18
<212> PRT
<213> *Caenorhabditis elegans*

<400> 271
Lys Thr His Ser Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr
1 5 10 15
Ala Lys

<210> 272
<211> 8
<212> PRT
<213> *Caenorhabditis elegans*

<400> 272
Lys Tyr Tyr Val Ser Pro Tyr Lys
1 5

<210> 273
<211> 18
<212> PRT
<213> *Caenorhabditis elegans*

<400> 273
Lys Phe Thr Ala His Pro Tyr Tyr Val Ser Arg Thr Pro Pro Arg Tyr
1 5 10 15
His Lys

<210> 274
<211> 67
<212> PRT
<213> *Caenorhabditis elegans*

<400> 274
Val Ile Lys Asn Leu Gly Tyr Met Val Asp Asn His Gly Phe Val Pro
1 5 10 15
Asn Gly Gly Arg Val Tyr Tyr Leu Thr Arg Ser Gln Pro Pro Leu Leu
20 25 30
Thr Pro Met Val Tyr Glu Tyr Tyr Met Ser Thr Gly Asp Leu Asp Phe
35 40 45
Val Met Glu Ile Leu Pro Thr Leu Asp Lys Glu Tyr Glu Phe Trp Ile
50 55 60
Lys Asn Arg
65

<210> 275
<211> 43
<212> PRT
<213> *Caenorhabditis elegans*

<400> 275
Ile Asn Leu Met Val Asp Gly Phe Val Pro Asn Gly Gly Arg Val Tyr
1 5 10 15
Tyr Leu Arg Ser Gln Pro Pro Leu Met Val Tyr Glu Tyr Thr Asp Phe
20 25 30
Val Glu Leu Pro Thr Leu Lys Glu Phe Trp Arg
35 40

<210> 276
<211> 67
<212> PRT
<213> *Caenorhabditis elegans*

<400> 276
Met Ile Arg Asn Leu Ala Ser Met Val Asp Lys Tyr Gly Phe Val Pro
1 5 10 15
Asn Gly Gly Arg Val Tyr Tyr Leu Gln Arg Ser Gln Pro Pro Phe Leu
20 25 30
Ala Ala Met Val Tyr Glu Leu Tyr Glu Ala Thr Asn Asp Lys Ala Phe
35 40 45
Val Ala Glu Leu Leu Pro Thr Leu Leu Lys Glu Leu Asn Phe Trp Asn
50 55 60
Glu Lys Arg

<210> 277
 <211> 84
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 277
 Ile Ile Pro Ala Asp Leu Asn Ala Phe Met Cys Ala Asn Ala Arg Ile
 1 5 10 15
 Leu Ala Ser Leu Tyr Glu Ile Ala Gly Asp Phe Lys Lys Val Lys Val
 20 25 30
 Phe Glu Gln Arg Tyr Thr Trp Ala Lys Arg Glu Met Arg Glu Leu His
 35 40 45
 Trp Asn Glu Thr Asp Gly Ile Trp Tyr Asp Tyr Asp Ile Glu Leu Lys
 50 55 60
 Thr His Ser Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr Ala
 65 70 75 80
 Lys Cys Tyr Asp

<210> 278
 <211> 31
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 278
 Pro Asp Leu Asn Cys Asn Ile Leu Tyr Glu Gly Asp Lys Phe Asn Thr
 1 5 10 15
 Asp Gly Trp Tyr Asp Tyr His Tyr Ser Ala Val Pro Leu Cys Tyr
 20 25 30

<210> 279
 <211> 84
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 279
 Val Leu Pro Val Asp Leu Asn Gly Leu Leu Cys Trp Asn Met Asp Ile
 1 5 10 15
 Met Glu Tyr Leu Tyr Glu Gln Ile Gly Asp Thr Lys Asn Ser Gln Ile
 20 25 30
 Phe Arg Asn Lys Arg Ala Asp Phe Arg Asp Thr Val Gln Asn Val Phe
 35 40 45
 Tyr Asn Arg Thr Asp Gly Thr Trp Tyr Asp Tyr Asn Leu Arg Thr Gln
 50 55 60
 Ser His Asn Pro Arg Phe Tyr Thr Ser Thr Ala Val Pro Leu Phe Thr
 65 70 75 80
 Asn Cys Tyr Asn

<210> 280
 <211> 48
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 280
Tyr Leu Glu Arg Gln Gly Leu Leu Lys Tyr Thr Lys Gly Leu Pro Thr
1 5 10 15
Ser Leu Ala Met Ser Ser Thr Gln Gln Trp Asp Lys Glu Asn Ala Trp
20 25 30
Pro Pro Met Ile His Met Val Ile Glu Gly Phe Arg Thr Thr Gly Asp
35 40 45

<210> 281
<211> 20
<212> PRT
<213> *Caenorhabditis elegans*

<400> 281
Gly Tyr Gly Pro Thr Ser Ser Gln Gln Trp Asp Asn Trp Pro His Met
1 5 10 15
Ile Glu Gly Arg
20

<210> 282
<211> 48
<212> PRT
<213> *Caenorhabditis elegans*

<400> 282
Phe Phe Gln Lys Met Gly Val Phe Thr Tyr Pro Gly Gly Ile Pro Thr
1 5 10 15
Ser Met Ser Gln Glu Ser Asp Gln Gln Trp Asp Phe Pro Asn Gly Trp
20 25 30
Ser Pro Asn Asn His Met Ile Ile Glu Gly Leu Arg Lys Ser Ala Asn
35 40 45

<210> 283
<211> 18
<212> PRT
<213> *Caenorhabditis elegans*

<400> 283
Glu Ile Ala Ser Ala Ala Glu Thr Gly Trp Asp Phe Ser Thr Arg Trp
1 5 10 15
Phe Ser

<210> 284
<211> 15
<212> PRT
<213> *Caenorhabditis elegans*

<400> 284
Ala Ser Ala Ala Glu Gly Trp Asp Phe Ser Thr Arg Trp Phe Ser
1 5 10 15

<210> 285
<211> 18

<212> PRT
<213> *Caenorhabditis elegans*

<400> 285
Asp Leu Ala Ser Ala Ala Glu Ser Gly Trp Asp Phe Ser Thr Arg Trp
1 5 10 15
Phe Ser

<210> 286
<211> 40
<212> PRT
<213> *Caenorhabditis elegans*

<400> 286
Lys Gln Phe Pro Tyr Tyr Gln Tyr Lys Ala Lys Leu Lys Val Pro Arg
1 5 10 15
Pro Glu Ser Tyr Arg Glu Asp Ser Glu Leu Ala Glu His Leu Gln Thr
20 25 30
Glu Ala Glu Lys Ile Gln Met Trp
35 40

<210> 287
<211> 18
<212> PRT
<213> *Caenorhabditis elegans*

<400> 287
Lys Phe Tyr Gln Tyr Lys Val Pro Arg Pro Glu Ser Tyr Arg Asp Leu
1 5 10 15
Ala Gln

<210> 288
<211> 40
<212> PRT
<213> *Caenorhabditis elegans*

<400> 288
Lys Ser Phe Lys Val Tyr Gln Tyr Lys Thr Ala Ser Asn Val Pro Arg
1 5 10 15
Pro Glu Ser Tyr Arg Val Asp Thr Gln Asn Ser Ala Lys Leu Ala Asn
20 25 30
Gly Ala Asp Gln Gln Gln Phe Tyr
35 40

<210> 289
<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 289
Gln Thr Gly Phe Gly Trp Thr Asn Gly Val Ile Leu Asp Leu Leu Asp
1 5 10 15
Lys Tyr Gly Asp Gln

<210> 290
<211> 14
<212> PRT
<213> *Caenorhabditis elegans*

<400> 290
Gln Gly Phe Gly Trp Asn Gly Ile Leu Asp Leu Leu Tyr Asp
1 5 10

<210> 291
<211> 21
<212> PRT
<213> *Caenorhabditis elegans*

<400> 291
Gln Asp Gly Phe Gly Trp Ser Asn Gly Ala Ile Leu Asp Leu Leu Leu
1 5 10 15
Thr Tyr Asn Asp Arg
20

<210> 292
<211> 27
<212> PRT
<213> *Caenorhabditis elegans*

<400> 292
Tyr Gly Asp Gln Phe Ala Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe
1 5 10 15
Ser Leu Ser Asn Ile Thr Phe Val Val Phe Ile
20 25

<210> 293
<211> 11
<212> PRT
<213> *Caenorhabditis elegans*

<400> 293
Tyr Phe Ala Ser Ser Ala Ser Phe Ser Phe
1 5 10

<210> 294
<211> 26
<212> PRT
<213> *Caenorhabditis elegans*

<400> 294
Tyr Asn Pro Phe Ala Ser Ser Ser Asp Ala Ser Ser Cys Pro Phe Ser
1 5 10 15
Thr Asn Ser Val Ile Phe Ser Ile Leu Val
20 25

<210> 295
<211> 9
<212> PRT
<213> *Caenorhabditis elegans*

<400> 295
Gly Gly Gly Gly Glu Tyr Glu Val Gln
1 5

<210> 296
<211> 7
<212> PRT
<213> *Caenorhabditis elegans*

<400> 296
Gly Gly Gly Glu Tyr Val Gln
1 5

<210> 297
<211> 9
<212> PRT
<213> *Caenorhabditis elegans*

<400> 297
Gly Ser Gly Gly Glu Tyr Asp Val Gln
1 5

<210> 298
<211> 14
<212> PRT
<213> *Caenorhabditis elegans*

<400> 298
Asn Gln Tyr Tyr Val Ser Asn Ala Val Pro Leu Tyr Ala Lys
1 5 10

<210> 299
<211> 7
<212> PRT
<213> *Caenorhabditis elegans*

<400> 299
Asn Tyr Tyr Val Leu Tyr Lys
1 5

<210> 300
<211> 14
<212> PRT
<213> *Caenorhabditis elegans*

<400> 300
Asn His Tyr Tyr Ile Ile Gln Met Val Ser Leu Tyr Thr Lys
1 5 10

<210> 301
<211> 30
<212> PRT
<213> *Caenorhabditis elegans*

<400> 301
Asp Gln Phe Ala Ser Ser Ser Thr Ala Ser Lys Phe Ser Phe Ser Leu
1 5 10 15
Ser Asn Ile Thr Phe Val Val Phe Ile Leu Tyr Ile Phe Ser
20 25 30

<210> 302
<211> 11
<212> PRT
<213> *Caenorhabditis elegans*

<400> 302
Asp Gln Phe Ser Ser Lys Phe Ser Phe Phe Ser
1 5 10

<210> 303
<211> 30
<212> PRT
<213> *Caenorhabditis elegans*

<400> 303
Asp Gln Phe Val Ile Ser Phe Ile Cys Ser Lys Phe Ser Ser Lys Asn
1 5 10 15
Lys Lys Leu Tyr Phe Cys Pro Ser His Phe Ser Leu Phe Ser
20 25 30

<210> 304
<211> 9
<212> PRT
<213> *Caenorhabditis elegans*

<220>
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<223> Xaa = Any Amino Acid

<400> 304
Gly Trp Asp Xaa Xaa Ile Ala Pro Lys
1 5

<210> 305
<211> 62
<212> PRT
<213> *Mus musculus*

<400> 305
Leu Cys Lys Glu Gly Ile Ser Asp Gly Ala Thr Met Lys Thr Phe Cys
1 5 10 15
Gly Thr Pro Glu Tyr Leu Ala Pro Glu Val Leu Glu Asp Asn Asp Tyr
20 25 30

Gly Arg Ala Val Asp Trp Trp Gly Leu Gly Val Val Met Tyr Glu Met
35 40 45
Met Cys Gly Arg Leu Pro Phe Tyr Asn Gln Asp His Glu Arg
50 55 60

<210> 306
<211> 9
<212> PRT
<213> *Caenorhabditis elegans*

<400> 306
Gln Ala Leu Thr Gln Met Asn Pro Lys
1 5

<210> 307
<211> 11
<212> PRT
<213> *Caenorhabditis elegans*

<400> 307
Gln Ala Leu Thr Gln Cys Val Asp Ser Met Arg
1 5 10

<210> 308
<211> 248
<212> PRT
<213> *Homo sapiens*

<400> 308
Ile Phe Arg Thr Ala Val Ser Ser Asn Arg Cys Arg Thr Glu Tyr Gln
1 5 10 15
Asn Ile Asp Leu Asp Cys Ala Tyr Ile Thr Asp Arg Ile Ile Ala Ile
20 25 30
Gly Tyr Pro Ala Thr Gly Ile Glu Ala Asn Phe Arg Asn Ser Lys Val
35 40 45
Gln Thr Gln Gln Phe Leu Thr Arg Arg His Gly Lys Gly Asn Val Lys
50 55 60
Val Phe Asn Leu Arg Gly Gly Tyr Tyr Asp Ala Asp Asn Phe Asp
65 70 75 80
Gly Asn Val Ile Cys Phe Asp Met Thr Asp His His Pro Pro Ser Leu
85 90 95
Glu Leu Met Ala Pro Phe Cys Arg Glu Ala Lys Glu Trp Leu Glu Ala
100 105 110
Asp Asp Lys His Val Ile Ala Val His Cys Lys Ala Gly Lys Gly Arg
115 120 125
Thr Gly Val Met Ile Cys Ala Leu Leu Ile Tyr Ile Asn Phe Tyr Pro
130 135 140
Ser Pro Arg Gln Ile Leu Asp Tyr Tyr Ser Ile Ile Thr Arg Lys Asn
145 150 155 160
Asn Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Ile Tyr Tyr Tyr
165 170 175
His Lys Leu Arg Glu Arg Glu Leu Asn Tyr Leu Pro Leu Arg Met Gln
180 185 190
Leu Ile Gly Val Tyr Val Glu Arg Pro Pro Lys Thr Trp Gly Gly Gly
195 200 205
Ser Lys Ile Lys Val Glu Val Gly Asn Gly Ser Thr Ile Leu Phe Lys

210	215	220
Pro Asp Pro Leu Ile Ile Ser Lys Ser Asn His Gln Arg Glu Arg Ala		
225	230	235
Thr Trp Leu Asn Asn Cys Asp Thr		
	245	

<210> 309
 <211> 249
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 309		
Ile Ile Lys Glu Ile Val Ser Arg Asn Lys Arg Arg Tyr Gln Glu Asp		
1 5 10 15		
Gly Phe Asp Leu Asp Leu Thr Tyr Ile Tyr Pro Asn Ile Ile Ala Met		
20 25 30		
Gly Phe Pro Ala Glu Arg Leu Glu Gly Val Tyr Arg Asn Asn Ile Asp		
35 40 45		
Asp Val Val Arg Phe Leu Asp Ser Lys His Lys Asn His Tyr Lys Ile		
50 55 60		
Tyr Asn Leu Cys Ala Glu Arg His Tyr Asp Thr Ala Lys Phe Asn Cys		
65 70 75 80		
Arg Val Ala Gln Tyr Pro Phe Glu Asp His Asn Pro Pro Gln Leu Glu		
85 90 95		
Leu Ile Lys Pro Phe Cys Glu Asp Leu Asp Gln Trp Leu Ser Glu Asp		
100 105 110		
Asp Asn His Val Ala Ala Ile His Cys Lys Ala Gly Lys Gly Arg Thr		
115 120 125		
Gly Val Met Ile Cys Ala Tyr Leu Leu His Arg Gly Lys Phe Leu Lys		
130 135 140		
Ala Gln Glu Ala Leu Asp Phe Tyr Gly Glu Val Arg Thr Arg Asp Lys		
145 150 155 160		
Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Val Tyr Tyr Ser		
165 170 175		
Tyr Leu Leu Lys Asn His Leu Asp Tyr Arg Pro Val Ala Leu Leu Phe		
180 185 190		
His Lys Met Met Phe Glu Thr Ile Pro Met Phe Ser Gly Gly Thr Cys		
195 200 205		
Asn Pro Gln Phe Val Val Cys Gln Leu Lys Val Lys Ile Tyr Ser Ser		
210 215 220		
Asn Ser Gly Pro Thr Arg Arg Glu Asp Lys Phe Asn Tyr Phe Glu Phe		
225 230 235 240		
Pro Gln Pro Leu Pro Val Cys Gly Asp		
245		

<210> 310
 <211> 962
 <212> PRT
 <213> *Caenorhabditis elegans*

<400> 310		
Met Val Thr Pro Pro Pro Asp Val Pro Ser Thr Ser Thr Arg Ser Met		
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Ala Arg Asp Leu Gln Glu Asn Pro Asn Arg Gln Pro Gly Glu Pro Arg		
20 25 30		
Val Ser Glu Pro Tyr His Asn Ser Ile Val Glu Arg Ile Arg His Ile		
35 40 45		

Phe Arg Thr Ala Val Ser Ser Asn Arg Cys Arg Thr Glu Tyr Gln Asn
 50 55 60
 Ile Asp Leu Asp Cys Ala Tyr Ile Thr Asp Arg Ile Ile Ala Ile Gly
 65 70 75 80
 Tyr Pro Ala Thr Gly Ile Glu Ala Asn Phe Arg Asn Ser Lys Val Gln
 85 90 95
 Thr Gln Gln Phe Leu Thr Arg Arg His Gly Lys Gly Asn Val Lys Val
 100 105 110
 Phe Asn Leu Arg Gly Gly Tyr Tyr Asp Ala Asp Asn Phe Asp Gly
 115 120 125
 Asn Val Ile Cys Phe Asp Met Thr Asp His His Pro Pro Ser Leu Glu
 130 135 140
 Leu Met Ala Pro Phe Cys Arg Glu Ala Lys Glu Trp Leu Glu Ala Asp
 145 150 155 160
 Asp Lys His Val Ile Ala Val His Cys Lys Ala Gly Lys Gly Arg Thr
 165 170 175
 Gly Val Met Ile Cys Ala Leu Leu Ile Tyr Ile Asn Phe Tyr Pro Ser
 180 185 190
 Pro Arg Gln Ile Leu Asp Tyr Tyr Ser Ile Ile Arg Thr Lys Asn Asn
 195 200 205
 Lys Gly Val Thr Ile Pro Ser Gln Arg Arg Tyr Ile Tyr Tyr Tyr His
 210 215 220
 Lys Leu Arg Glu Arg Glu Leu Asn Tyr Leu Pro Leu Arg Met Gln Leu
 225 230 235 240
 Ile Gly Val Tyr Val Glu Arg Pro Pro Lys Thr Trp Gly Gly Ser
 245 250 255
 Lys Ile Lys Val Glu Val Gly Asn Gly Ser Thr Ile Leu Phe Lys Pro
 260 265 270
 Asp Pro Leu Ile Ile Ser Lys Ser Asn His Gln Arg Glu Arg Ala Thr
 275 280 285
 Trp Leu Asn Asn Cys Asp Thr Pro Asn Glu Phe Asp Thr Gly Glu Gln
 290 295 300
 Lys Tyr His Gly Phe Val Ser Lys Arg Ala Tyr Cys Phe Met Val Pro
 305 310 315 320
 Glu Asp Ala Pro Val Phe Val Glu Gly Asp Val Arg Ile Asp Ile Arg
 325 330 335
 Glu Ile Gly Phe Leu Lys Lys Phe Ser Asp Gly Lys Ile Gly His Val
 340 345 350
 Trp Phe Asn Thr Met Phe Ala Cys Asp Gly Gly Leu Asn Gly Gly His
 355 360 365
 Phe Glu Tyr Val Asp Lys Thr Gln Pro Tyr Ile Gly Asp Asp Thr Ser
 370 375 380
 Ile Gly Arg Lys Asn Gly Met Arg Arg Asn Glu Thr Pro Met Arg Lys
 385 390 395 400
 Ile Asp Pro Glu Thr Gly Asn Glu Phe Glu Ser Pro Trp Gln Ile Val
 405 410 415
 Asn Pro Pro Gly Leu Glu Lys His Ile Thr Glu Glu Gln Ala Met Glu
 420 425 430
 Asn Tyr Thr Asn Tyr Gly Met Ile Pro Pro Arg Tyr Thr Ile Ser Lys
 435 440 445
 Ile Leu His Glu Lys His Glu Lys Gly Ile Val Lys Asp Asp Tyr Asn
 450 455 460
 Asp Arg Lys Leu Pro Met Gly Asp Lys Ser Tyr Thr Glu Ser Gly Lys
 465 470 475 480
 Ser Gly Asp Ile Arg Gly Val Gly Gly Pro Phe Glu Ile Pro Tyr Lys
 485 490 495
 Ala Glu Glu His Val Leu Thr Phe Pro Val Tyr Glu Met Asp Arg Ala
 500 505 510
 Leu Lys Ser Lys Asp Leu Asn Asn Gly Met Lys Leu His Val Val Leu

515	520	525	
Arg Cys Val Asp Thr Arg Asp	Ser Lys Met Met	Glu Lys Ser Glu Val	
530	535	540	
Phe Gly Asn Leu Ala Phe His Asn Glu Ser	Thr Arg Arg	Leu Gln Ala	
545	550	555	560
Leu Thr Gln Met Asn Pro Lys Trp Arg	Pro Glu Pro Cys Ala	Phe Gly	
565	570	575	
Ser Lys Gly Ala Glu Met His Tyr	Pro Pro Ser Val Arg	Tyr Ser Ser	
580	585	590	
Asn Asp Gly Lys Tyr Asn Gly	Ala Cys Ser Glu Asn	Leu Val Ser Asp	
595	600	605	
Phe Phe Glu His Arg Asn Ile Ala Val	Leu Asn Arg	Tyr Cys Arg Tyr	
610	615	620	
Phe Tyr Lys Gln Arg Ser Thr Ser Arg	Arg Tyr Pro Arg	Lys Phe	
625	630	635	640
Arg Tyr Cys Pro Leu Ile Lys Lys His	Phe Tyr Ile Pro Ala Asp	Thr	
645	650	655	
Asp Asp Val Asp Glu Asn Gly Gln	Pro Phe His Ser	Pro Glu His	
660	665	670	
Tyr Ile Lys Glu Gln Glu Lys Ile Asp	Ala Glu Lys Ala	Ala Lys Gly	
675	680	685	
Ile Glu Asn Thr Gly Pro Ser Thr Ser Gly	Ser Ala Pro Gly	Thr	
690	695	700	
Ile Lys Lys Thr Glu Ala Ser Gln Ser Asp	Lys Val Lys Pro Ala	Thr	
705	710	715	720
Glu Asp Glu Leu Pro Pro Ala Arg Leu	Pro Asp Asn Val Arg	Arg Phe	
725	730	735	
Pro Val Val Gly Val Asp Phe Glu Asn	Pro Glu Glu Glu	Ser Cys Glu	
740	745	750	
His Lys Thr Val Glu Ser Ile Ala Gly	Phe Glu Pro Leu	Glu His Leu	
755	760	765	
Phe His Glu Ser Tyr His Pro Asn Thr Ala	Gly Asn Met Leu Arg	Gln	
770	775	780	
Asp Tyr His Thr Asp Ser Glu Val Lys	Ile Ala Glu Gln Glu Ala	Lys	
785	790	795	800
Ala Phe Val Asp Gln Leu Leu Asn Gly	Gln Gly Val Leu Gln Glu	Phe	
805	810	815	
Met Lys Gln Phe Lys Val Pro Ser Asp	Asn Ser Phe Ala Asp	Tyr Val	
820	825	830	
Thr Gly Gln Ala Glu Val Phe Lys	Ala Gln Ile Ala	Leu Leu Glu Gln	
835	840	845	
Ser Glu Asp Phe Gln Arg Val Gln Ala	Asn Ala Glu Glu Val Asp	Leu	
850	855	860	
Glu His Thr Leu Gly Glu Ala Phe Glu	Arg Phe Gly His Val	Val Glu	
865	870	875	880
Glu Ser Asn Gly Ser Ser Lys Asn Pro	Lys Ala Leu Lys Thr Arg	Glu	
885	890	895	
Gln Met Val Lys Glu Thr Gly Lys Asp	Thr Gln Lys Thr Arg	Asn His	
900	905	910	
Val Leu Leu His Leu Glu Ala Asn His	Arg Val Gln Ile	Glu Arg Arg	
915	920	925	
Glu Thr Cys Pro Glu Leu His Pro Glu	Asp Lys Ile Pro Arg	Ile Ala	
930	935	940	
His Phe Ser Glu Asn Ser Phe Ser Asp	Ser Asn Phe Asp	Gln Ala Ile	
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Tyr Leu			

<210> 311
 <211> 3304
 <212> DNA
 <213> *Caenorhabditis elegans*

<400> 311

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tgtatcttcc	aatcggtgtc	gcaccgagta	ccaaaatatc	gacctagatt	gtgcatatat	240
cacagaccga	atcatagcta	tcggttatcc	agcaacagga	atcgaagcga	atttccgtaa	300
ctcaaaaagt	caaactcaac	aatttctgac	caggcggcac	ggaaaggc	acgtgaaggt	360
gtttaacctg	cgcggtg	actactacga	tgcggataac	ttcgatggaa	atgttattt	420
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aattccatca	caacgacgct	acatttacta	ctaccataag	cttcgtgaac	gtgagctcaa	720
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ctgtgatacg	cctaacgaat	tcgacaccgg	agagcaaaaa	tatcatggat	ttgtttccaa	960
gagagcatac	tgtttatgg	tgccagaaga	tgctccagta	tttgcgaag	gagatgtcg	1020
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ttggttcaat	acaatgttcg	catgtgatgg	aggactcaac	ggtggacatt	tcgagta	1140
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gtggcaaata	gtgaatc	ctggactg	aaaacatatt	acggaggaac	aagcaatgga	1320
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caaatcatac	acgaaatcag	gaaaaagtgg	agatattcga	ggagt	cggt	1500
gataccat	aaagctgagg	aacatgttct	cacatttcca	gtttatgaaa	tggatcgagc	1560
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caatgatgga	aagtataatg	gac	tgagaac	gttagc	tttgc	1860
cagaaatatt	gccgttctt	atcgatatt	ccgatattt	tacaagcaac	gcag	1920
tcgaagccgt	tatccaagaa	aattcagata	ctgtcctct	atcaagaa	atttctacat	1980
tccagctat	accatgtat	ttgatgaaa	tggcaacc	tttccact	caccagag	2040
ttacattaaa	gaacaggaaa	aaatagacgc	agagaaagca	gctaaaggaa	ttgaaaatac	2100
tggacccagt	acttcaggat	caagt	cgaaactatc	aagaaaacgg	aagctt	2160
atccgacaag	gtgaagccgg	caactgaaga	cgaactt	cctgcgaggc	taccggataa	2220
tgtgcgaaga	tttccagtcg	tcggcgtt	ttcgaaaat	ccggaaag	aatcgtgt	2280
acacaaaacc	gtagagtcaa	tagtgg	tgaaccact	gaacatct	tccatgaa	2340
ataccatcca	aatacggccg	gtaacatgt	gcgtcaggat	tatcacact	attcggaa	2400
aaaaatagct	gaacaagagg	caaaagc	cg	ttgctt	atg	2460
attacaagag	tttatgaa	aaatcaag	accatcg	gac	at	2520
aaccggacag	gccgaagtt	ttaaagcaca	gat	ctgg	g	2580
tcaacgagtt	caagcgaat	cagagga	cgat	tca	gt	2640
tgagcgattc	ggcacgtt	tagaaga	gaat	ttct	g	2700
aaaaactcga	gaacaaatgg	tgaaagaa	tggcaag	actc	cc	2760
tgtgcttcta	catttggaa	cta	tgt	gcaat	gt	2820
ggagctacat	ccagaggata	aaatccaa	aatt	gtc	ttt	2880
ggattcgaat	tttgc	at	ttt	ccgaaa	acag	2940
ttcttcttac	tgacc	ttt	ttt	aaactt	ttt	3000
gtatcattca	ctt	ttt	ttt	ttt	ttt	3060
tatattcata	ttat	at	ttt	ttt	ttt	3120
ggtgaaaaat	agcaatt	ttt	ttt	ttt	ttt	3180
tttcc	tttcc	tttcc	tttcc	tttcc	tttcc	

ttgttaattca cattgcgggt catcaactaat cctatggct ttaacacaat tctcccataa	3240
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<210> 312

<211> 1642

<212> DNA

<213> *Caenorhabditis elegans*

<400> 312

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gatggcatcc gcaatgaagt ttcaataacta ctcgaagaaa gctgctggaa agacaatgtc	120
taatagtgtc tccatgtcca gtgacaatcg catggaggat tttaaacgtc gtttgcgtcg	180
aagtggatcg ttaggaattc catttgtccc agaagaagat gttaaacaac tcttcacacc	240
aactcgtact gttcgtcgag aagcatctat tcgcgaaggg gatgaggaag aaggagtaca	300
aattctcaca ataattgtca agtcaagtgc tggatcgag gatatctcaa aaatgattgc	360
aaacacctccct gatcacactc gtatcaaaca ttggagact cgtgacagtc aagatgaaag	420
ttccaaaact atggatgttc ttcttagat ttagctctt cattatggaa aacaagaagc	480
aatggatctt atgagactta atgggcttga tggatcgag gatgacgacaa ccggttctga	540
aactgcaata aaagagcaat atacagagcc tggatctgat gatgcgacaa ccggttctga	600
atggtttcca aaaagtattt atgatttggaa tatttgcataaaaagatgatgatgtatgg	660
agcagggctg gacgctgatc atcctgggtt caaagatacc gatgatcgatc aacgtcgaat	720
gatgtttgcgaaacttggcgtc tcaattacaa acacggtagg ccaattccgc gaaccgaata	780
tacatcatcc gaacggaaaaa cttggggaaat tatataaaaaaattgagag aattgcacaa	840
aaagcacgca tgcaaggactt ttcttgataa cttggatcttgcataaaaactgg	900
ctcgaaaaat aatattccgc aactagaaga tatctgcaag ttttgcataaaaactgg	960
attccgtgtt cgcccagtcg ccggataactt atcagctcgatgattcttgcgatgtttgc	1020
atatcggtc ttcttctgca ctcaataacgt tcgcatcat ggcgatccat ttacactcc	1080
agaaccagac accgttcacg agctcatggg tcacatggct ctattcgatc atccagat	1140
tgctcagttt tctcaagaga ttggattagc ttctcttgcataaaaactgg	1200
gaagcttgca acactctact tctttccat tgaatttgcataaaaactgg	1260
cgattctcca gtaaaagaaa atggatcaaa tcatgaaaga ttggatcgatgtttgc	1320
acttctgagc agtgcggcg agttgcaaca tgccgttgcataaaaactgg	1380
ttttgatccg gatgtgttgcataaaaactgg	1440
tttctataact agaaatttttgcataaaaactgg	1500
gaaacgtccc ttcattgttc gttacaaccc atacacagaa agcgtcgaag ttctcaacaa	1560
ctcccggttcc attatgttgcataaaaactgg	1620
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<210> 313

<211> 532

<212> PRT

<213> *Caenorhabditis elegans*

<400> 313

Met Asp Ser Leu Phe Gln Met Ala Ser Ala Met Lys Phe Gln Tyr Tyr	
1 5 10 15	
Ser Lys Lys Ala Ala Gly Lys Thr Met Ser Asn Ser Val Ser Met Ser	
20 25 30	
Ser Asp Asn Arg Met Glu Asp Phe Lys Arg Arg Phe Arg Arg Ser Gly	
35 40 45	
Ser Leu Gly Ile Pro Phe Val Pro Glu Glu Asp Val Lys Gln Leu Phe	
50 55 60	
Thr Pro Thr Arg Thr Val Arg Arg Glu Ala Ser Ile Arg Glu Gly Asp	
65 70 75 80	
Glu Glu Glu Gly Val Gln Ile Leu Thr Ile Ile Val Lys Ser Ser Arg	
85 90 95	
Val Ser Glu Asp Ile Ser Lys Met Ile Ala Asn Leu Pro Asp His Thr	
100 105 110	

Arg Ile Lys His Leu Glu Thr Arg Asp Ser Gln Asp Gly Ser Ser Lys
 115 120 125
 Thr Met Asp Val Leu Leu Glu Ile Glu Leu Phe His Tyr Gly Lys Gln
 130 135 140
 Glu Ala Met Asp Leu Met Arg Leu Asn Gly Leu Asp Val His Glu Val
 145 150 155 160
 Ser Ser Thr Ile Arg Pro Thr Ala Ile Lys Glu Gln Tyr Thr Glu Pro
 165 170 175
 Gly Ser Asp Asp Ala Thr Thr Gly Ser Glu Trp Phe Pro Lys Ser Ile
 180 185 190
 Tyr Asp Leu Asp Ile Cys Ala Lys Arg Val Ile Met Tyr Gly Ala Gly
 195 200 205
 Leu Asp Ala Asp His Pro Gly Phe Lys Asp Thr Glu Tyr Arg Gln Arg
 210 215 220
 Arg Met Met Phe Ala Glu Leu Ala Leu Asn Tyr Lys His Gly Glu Pro
 225 230 235 240
 Ile Pro Arg Thr Glu Tyr Thr Ser Ser Glu Arg Lys Thr Trp Gly Ile
 245 250 255
 Ile Tyr Arg Lys Leu Arg Glu Leu His Lys Lys His Ala Cys Lys Gln
 260 265 270
 Phe Leu Asp Asn Phe Glu Leu Leu Glu Arg His Cys Gly Tyr Ser Glu
 275 280 285
 Asn Asn Ile Pro Gln Leu Glu Asp Ile Cys Lys Phe Leu Lys Ala Lys
 290 295 300
 Thr Gly Phe Arg Val Arg Pro Val Ala Gly Tyr Leu Ser Ala Arg Asp
 305 310 315 320
 Phe Leu Ala Gly Leu Ala Tyr Arg Val Phe Phe Cys Thr Gln Tyr Val
 325 330 335
 Arg His His Ala Asp Pro Phe Tyr Thr Pro Glu Pro Asp Thr Val His
 340 345 350
 Glu Leu Met Gly His Met Ala Leu Phe Ala Asp Pro Asp Phe Ala Gln
 355 360 365
 Phe Ser Gln Glu Ile Gly Leu Ala Ser Leu Gly Ala Ser Glu Glu Asp
 370 375 380
 Leu Lys Lys Leu Ala Thr Leu Tyr Phe Phe Ser Ile Glu Phe Gly Leu
 385 390 395 400
 Ser Ser Asp Asp Ala Ala Asp Ser Pro Val Lys Glu Asn Gly Ser Asn
 405 410 415
 His Glu Arg Phe Lys Val Tyr Gly Ala Gly Leu Leu Ser Ser Ala Gly
 420 425 430
 Glu Leu Gln His Ala Val Glu Gly Ser Ala Thr Ile Ile Arg Phe Asp
 435 440 445
 Pro Asp Arg Val Val Glu Gln Glu Cys Leu Ile Thr Thr Phe Gln Ser
 450 455 460
 Ala Tyr Phe Tyr Thr Arg Asn Phe Glu Glu Ala Gln Gln Lys Leu Arg
 465 470 475 480
 Met Phe Thr Asn Asn Met Lys Arg Pro Phe Ile Val Arg Tyr Asn Pro
 485 490 495
 Tyr Thr Glu Ser Val Glu Val Leu Asn Asn Ser Arg Ser Ile Met Leu
 500 505 510
 Ala Val Asn Ser Leu Arg Ser Asp Ile Asn Leu Leu Ala Gly Ala Leu
 515 520 525
 His Tyr Ile Leu
 530

<210> 314
 <211> 817
 <212> DNA

<213> *Caenorhabditis elegans*

<400> 314

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agtgtcaaaa	actggattcc	gtgttcgccc	agtgcggga	tacttacag	ctcgatgtt	180
cttggcaggt	cttgcata	gtgtcttctt	ctgactcaa	tacgttcg	atcatgccga	240
tccattttac	actccagaac	cagacaccgt	tcacgagtc	atgggtcaca	tggctctatt	300
cgctgatcca	gatttgctc	agtttctca	agagattgga	ttagcttctc	ttggagcatc	360
agaggaagat	ttgaagaagc	ttgcaacact	ctacttctt	tccattgaat	ttggctctc	420
gtctgatgac	gctgccgatt	ctccagtaaa	agaaaaatgga	tcaaattatg	aaagattaa	480
agtatacgg	gcaggacttc	tgagcagtgc	tggcgagtt	caacatgccg	ttgaggtag	540
tgcaaccatt	attcgtttg	atccggatcg	tgttgtttag	caagaatgtc	tcattactac	600
tttccagtca	gcgtatttct	atactagaaa	tttgaagag	gcccagcaga	aactcagaat	660
gttcaccaac	aacatgaaac	gtcccttcat	tgttcgttac	aaccatatac	cagaaagcgt	720
cgaagttctc	aacaactccc	gttccattat	gttggcagt	aactctctcc	gctcagacat	780
caacctgctc	gccggagctc	tccactacat	cctgtag			817

<210> 315

<211> 45

<212> PRT

<213> *Caenorhabditis elegans*

<400> 315

Met	Asp	Ser	Leu	Phe	Gln	Met	Ala	Ser	Ala	Met	Lys	Phe	Gln	Tyr	Tyr
1			5			10					15				
Ser	Lys	Lys	Ala	Ala	Gly	Lys	Thr	Met	Ser	Asn	Ser	Val	Lys	Asn	Trp
			20				25					30			
Ile	Pro	Cys	Ser	Pro	Ser	Arg	Arg	Ile	Leu	Ile	Ser	Ser			
			35				40				45				

<210> 316

<211> 466

<212> DNA

<213> *Caenorhabditis elegans*

<400> 316

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gactttgagg	cttcggctg	ggactggatc	atcgcaccta	agcgctacaa	ggccaactac	180
tgctccggcc	agtgggagta	catgttcatg	caaaaatatc	cgcataccca	tgtggtgcag	240
caggccaaatc	caagaggta	tgctggccc	tgttgcaccc	ccaccaagat	gtccccaaatc	300
aacatgctct	acttcaatga	caagcagcag	attatctacg	gcaagatccc	tggcatggtg	360
gtggatcgct	gtggctgctc	ttaagggtgg	ggatagagga	tgcctcccc	acagaccgta	420
ccccaaagacc	catagccctg	cccaatccac	cgcctgatcc	aaacat		466

<210> 317

<211> 128

<212> PRT

<213> *Caenorhabditis elegans*

<400> 317

Ile	Arg	His	Glu	His	Gly	Ala	Ser	Ser	Pro	Arg	Glu	His	Lys	Thr	Phe
1			5			10					15				
Pro	Ala	Glu	Pro	Gly	Ser	Gly	Leu	Arg	Arg	Asp	Ser	Ser	Glu	Ser	Arg
			20				25				30				
Cys	Cys	Arg	Tyr	Pro	Leu	Thr	Val	Asp	Phe	Glu	Ala	Phe	Gly	Trp	Asp
			35				40			45					

Trp	Ile	Ile	Ala	Pro	Lys	Arg	Tyr	Lys	Ala	Asn	Tyr	Cys	Ser	Gly	Gln
50					55				60						
Trp	Glu	Tyr	Met	Phe	Met	Gln	Lys	Tyr	Pro	His	Thr	His	Leu	Val	Gln
65					70				75				80		
Gln	Ala	Asn	Pro	Arg	Gly	Tyr	Ala	Gly	Pro	Cys	Cys	Thr	Pro	Thr	Lys
					85				90				95		
Met	Ser	Pro	Ile	Asn	Met	Leu	Tyr	Phe	Asn	Asp	Lys	Gln	Gln	Ile	Ile
					100			105				110			
Tyr	Gly	Lys	Ile	Pro	Leu	Ala	Met	Val	Val	Asp	Arg	Cys	Gly	Cys	Ser
					115			120				125			

<210> 318
<211> 9
<212> DNA
<213> Homo sapiens

<400> 318
caaaaactaa

9

<210> 319
<211> 20
<212> DNA
<213> Caenorhabditis elegans

<400> 319
ccactatggc cgagatttcc

20

<210> 320
<211> 44
<212> DNA
<213> Caenorhabditis elegans

<400> 320
ccagtgaaaaa gttcttctcc ttttttcctc ttctcgaatt cgga

44

<210> 321
<211> 21
<212> DNA
<213> Caenorhabditis elegans

<400> 321
cttccttcttc tcgaattcgg c

21

<210> 322
<211> 8
<212> PRT
<213> Caenorhabditis elegans

<400> 322
Gly Arg Lys Gly Phe Pro His Val
1 5

<210> 323
<211> 7
<212> PRT
<213> Caenorhabditis elegans

<220>
<221> VARIANT
<222> (1)...(7)
<223> Xaa = Any Amino Acid

<400> 323
Arg Xaa Xaa Ile Xaa Xaa Gly
1 5

<210> 324
<211> 7
<212> PRT
<213> *Caenorhabditis elegans* or *Homo sapiens*

<400> 324
Cys Gly Cys Cys Cys Cys Cys
1 5

<210> 325
<211> 79
<212> PRT
<213> *Homo sapiens* or *Caenorhabditis elegans*

<400> 325
Val Leu Asp Asp Tyr Gly Arg Val Asp Trp Trp Gly Gly Val Val Met
1 5 10 15
Tyr Glu Met Met Cys Gly Arg Leu Pro Phe Tyr Asp His Lys Leu Phe
20 25 30
Glu Leu Ile Arg Phe Pro Leu Glu Ala Leu Leu Gly Leu Leu Lys Asp
35 40 45
Pro Thr Gln Arg Leu Gly Gly Glu Asp Ala Glu Ile Phe Phe Trp
50 55 60
Tyr Lys Pro Pro Lys Pro Val Ser Glu Thr Asp Thr Tyr Phe Asp
65 70 75

<210> 326
<211> 48
<212> PRT
<213> *Homo sapiens* or *Caenorhabditis elegans*

<400> 326
Thr Met Phe Leu Lys Leu Gly Lys Gly Thr Phe Gly Lys Val Ile Leu
1 5 10 15
Lys Glu Lys Thr Tyr Ala Lys Ile Leu Lys Lys Val Ile Ala Glu Val
20 25 30
Ala His Thr Leu Thr Glu Asn Arg Val Leu Gln His Pro Phe Leu Thr
35 40 45

<210> 327
<211> 27
<212> DNA
<213> *Caenorhabditis elegans*

<400> 327
caagcggttga ctcaaatgaa tccaaaa

<210> 328
<211> 55
<212> DNA
<213> *Caenorhabditis elegans*

<400> 328
caagcggtga ctcaatgcgt tgactcaatg cgttgactcg ttgacgaatc caaaa

55